

Causal models and how to refute them.

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How a short nap can raise the risk of diabetes: Study finds people who have a siesta are more likely to have high blood pressure and high cholesterol

- Napping for more than 30 minutes at a time can raise the risk of diabetes, according to a new study
- It can also increase likelihood of high blood pressure and high cholesterol

By PAT HAGAN

PUBLISHED: 01:04, 21 September 2013 | **UPDATED:** 10:34, 21 September 2013

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How a short nap can raise blood pressure and high cholesterol: Study finds people who take a siesta are more likely to have high blood pressure and high cholesterol

- Napping for more than 30 minutes at a time is associated with a higher risk of impaired fasting plasma glucose and diabetes mellitus in older adults: results from the Dongfeng–Tongji cohort of retired workers
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ELSEVIER

Sleep Medicine

Volume 14, Issue 10, October 2013, Pages 950–954



Original Article

Longer habitual afternoon napping is associated with a higher risk for impaired fasting plasma glucose and diabetes mellitus in older adults: results from the Dongfeng–Tongji cohort of retired workers

Weimin Fang^{a, b}, Zhongliang Li^a, Li Wu^a, Zhongqiang Cao^a, Yuan Liang^{a, c}, Handong Yang^d, Youjie Wang^{a, b, e, f}, Tangchun Wu^a

^aMinistry of Education Key Laboratory of Environment and Health, School of Public Health, Tongji Medical College, Huazhong University of Science and Technology, China

^bDepartment of Maternal and Child Health, School of Public Health, Tongji Medical College, Huazhong University of Science and Technology, China

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^dDongfeng General Hospital, Dongfeng Motor Corporation and Hubei University of Medicine, China

Abstract

Objectives

Afternoon napping is a common habit in China. We used data obtained from the Dongfeng–Tongji cohort to examine if duration of habitual afternoon napping was associated with risks for impaired fasting plasma glucose (IFG) and diabetes mellitus (DM) in a Chinese elderly population.

Methods

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“Dr Matthew Hobbs, head of research for Diabetes UK, said there was no proof that napping **actually caused** diabetes.”

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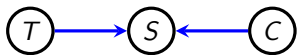
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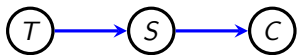
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This is the basis of some causal search algorithms (e.g. PC, FCI).

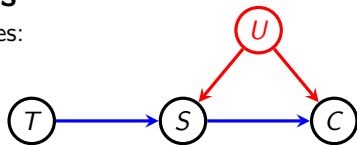
Note: other methods (e.g. integer programming) are also used.

In order to do this well, we need to understand in what ways causal models will be **observationally** different.

When everything is observed this is (mathematically) easy.

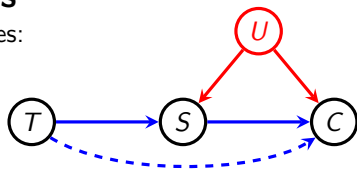
Hidden Variables

Instrumental variables:



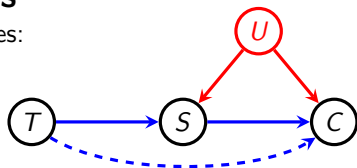
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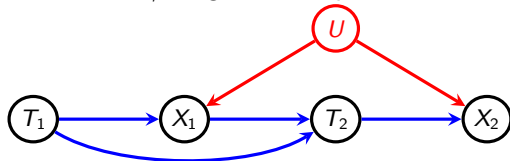


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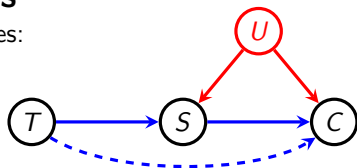


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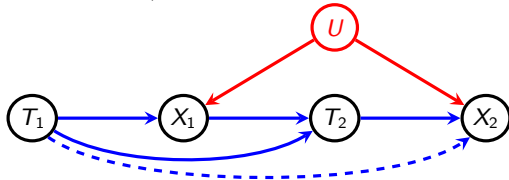


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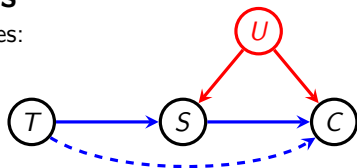


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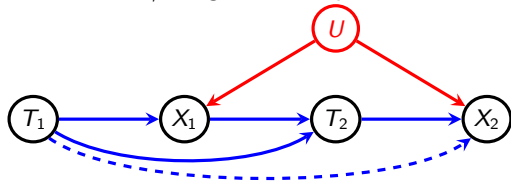


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Principal aims:

- be able to test causal models;
- identify and bound causal effects;
- use constraints for model search.

The Holy Grail: Structure Learning

Truth:

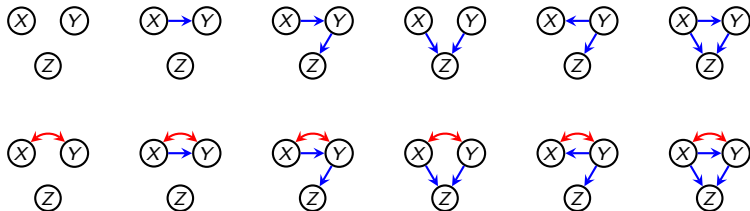


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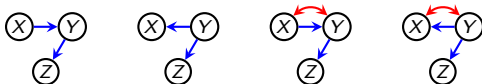
...return list of models which are compatible with data.

Experimental Design

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We could then identify an experiment to distinguish remaining models:

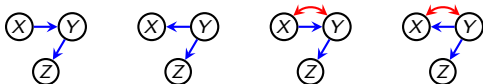


Experimental Design

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intervene on X:




...return list of models which are compatible with data.

To do this we need to know what constraints the model places on the distribution (the focus of this talk).

Outline


- 1 DAG Models
- 2 An Example
- 3 mDAGs
- 4 Inequalities
- 5 Testing, Fitting and Searching

Directed Acyclic Graphs

vertices 

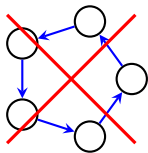
edges 

Directed Acyclic Graphs


vertices 

edges 

no directed cycles

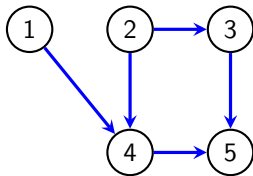
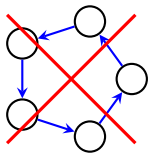


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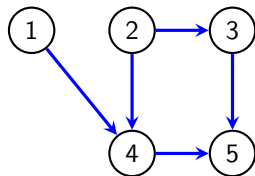
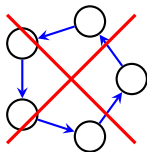
directed acyclic graph (DAG), \mathcal{G}

Directed Acyclic Graphs

vertices 

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directed acyclic graph (DAG), \mathcal{G}

If $w \rightarrow v$ then w is a **parent** of v : $\text{pa}_{\mathcal{G}}(4) = \{1, 2\}$.

If $w \rightarrow \dots \rightarrow v$ then w is a **ancestor** of v .

An **ancestral set** contains all its own ancestors.

DAG Models (aka Bayesian Networks)

vertex



random variable



X_a

DAG Models (aka Bayesian Networks)

vertex

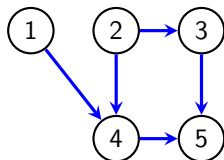


random variable

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graph \mathcal{G}



model $\mathcal{M}(\mathcal{G})$

$$p(x_1, \dots, x_k) = \prod_i p(x_i | x_{\text{pa}(i)}).$$

(factorization)



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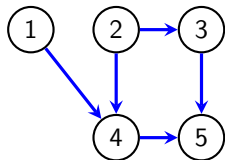


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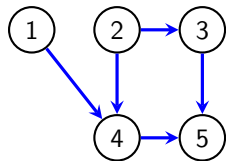


So in example above:

$$p(x_V) = p(x_1) \cdot p(x_2) \cdot p(x_3 | x_2) \cdot p(x_4 | x_1, x_2) \cdot p(x_5 | x_3, x_4)$$

DAG Models

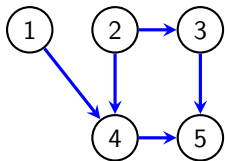
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pick a topological ordering
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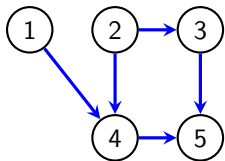
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Can *always* factorize a joint distribution as:

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The model is the same as setting

$$p(x_i | x_1, x_2, \dots, x_{i-1}) = p(x_i | x_{\text{pa}(i)}), \quad \text{for each } i.$$

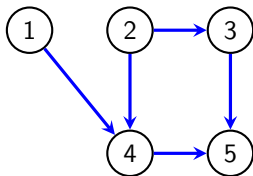
Thus $\mathcal{M}(\mathcal{G})$ is precisely distributions such that:

$$X_i \perp\!\!\!\perp X_{[i-1] \setminus \text{pa}(i)} \mid X_{\text{pa}(i)}, \quad i \in V.$$

This is a constraint-based perspective.

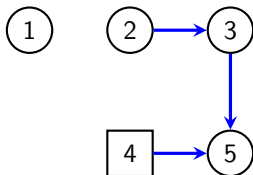
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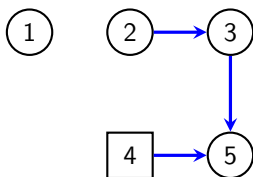
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All other terms preserved.

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Marginalization

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- over 10,000 Wisconsin high-school graduates from 1957;
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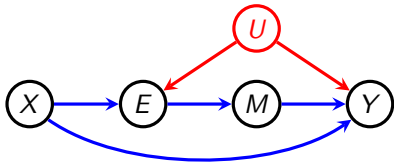
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U unmeasured confounding.



Marginalization

Note we don't want to make assumptions about U ; so this is **not** a latent variable model in the usual sense.

Model is defined (implicitly) by an integral:

$$p(x, e, m, y) = \int p(u) p(x) p(e | x, u) p(m | e) p(y | x, m, u) du$$

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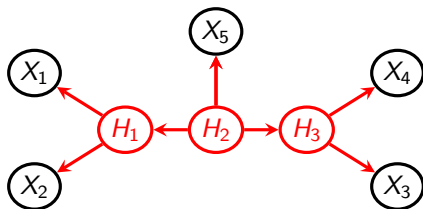
We aim to study the set of distributions constructed in this way.

Strategy: study constraints satisfied by these models.

Latent Variable Models

Traditional latent variable models would assume that the hidden variables are (e.g.) Gaussian, or discrete with some fixed number of states.

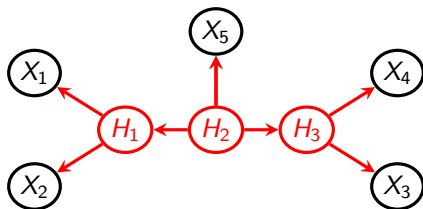
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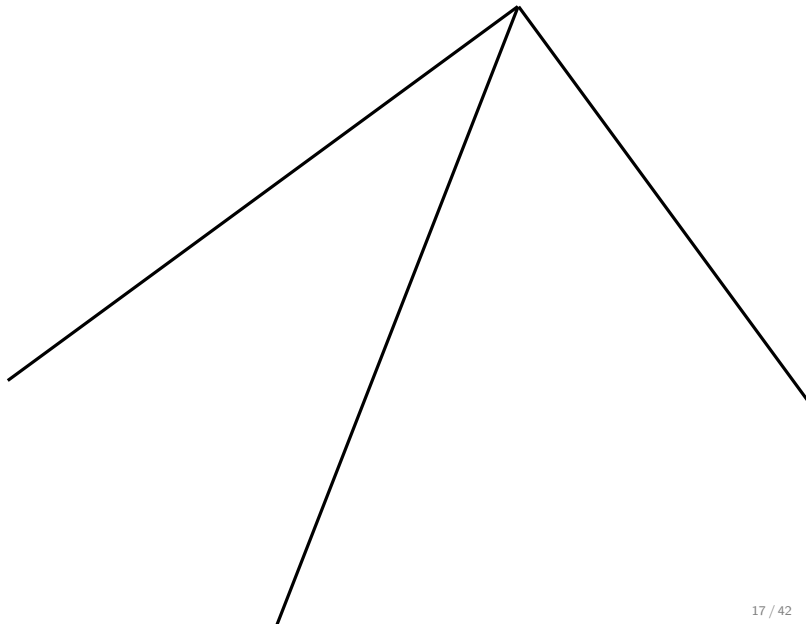
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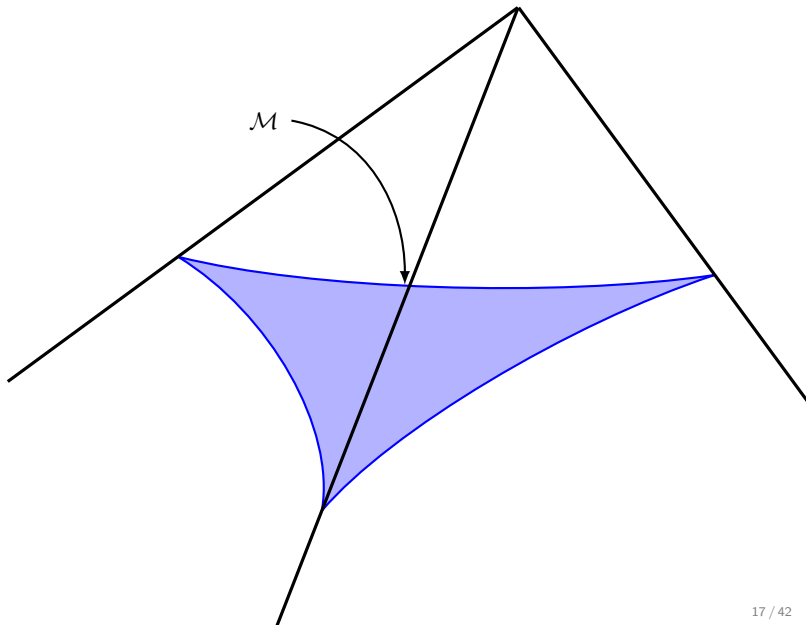
But:

- assumptions may be wrong!
- latent variables lead to singularities and nasty statistical properties (see e.g. Drton, Sturmfels and Sullivant, 2009)

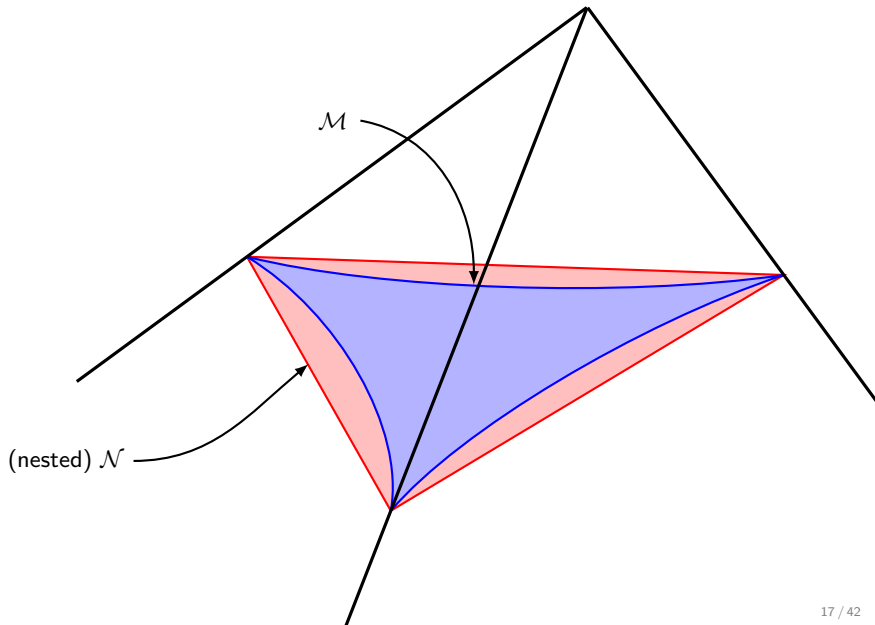
Getting the Picture



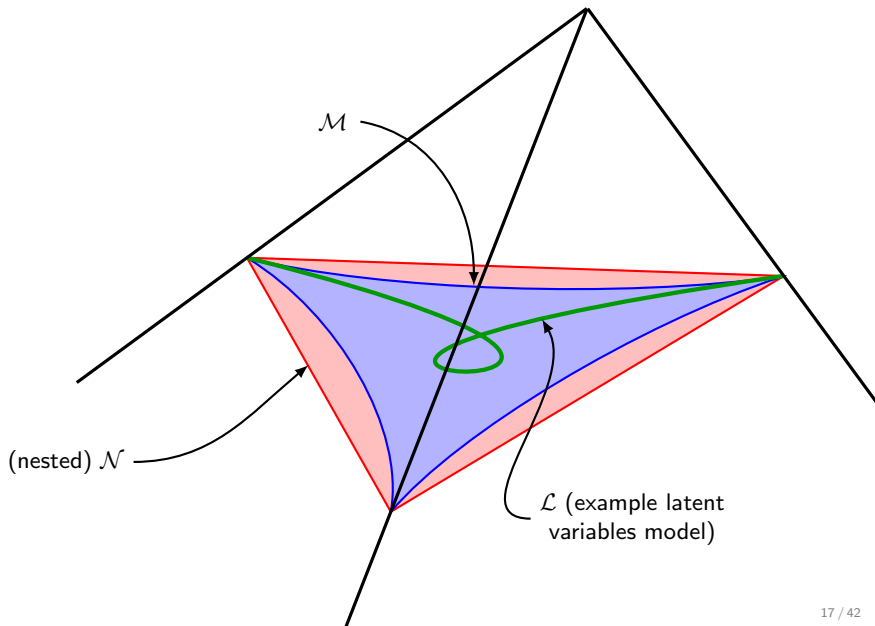
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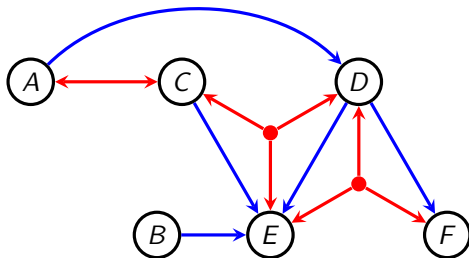


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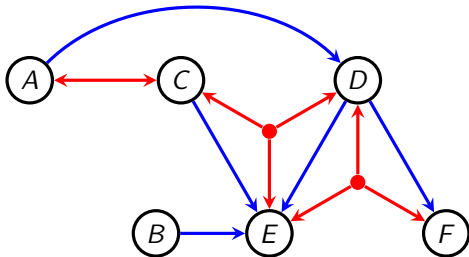
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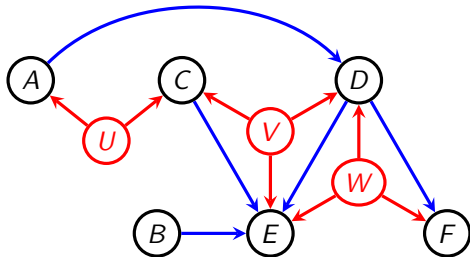
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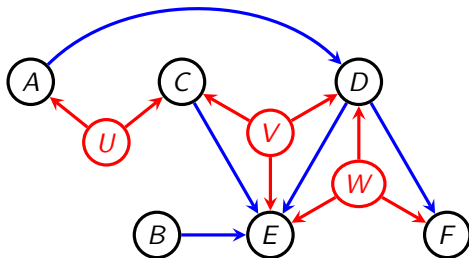


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Can put the latents back: call this the **canonical DAG** $\bar{\mathcal{G}}$.

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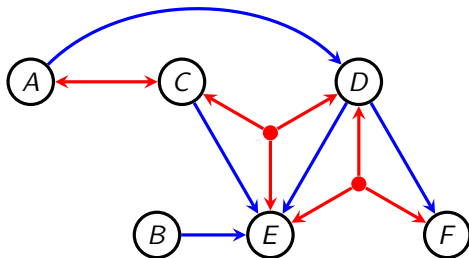
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The **marginal model** is denoted $\mathcal{M}(\mathcal{G})$.

Model Description

We can write down a causal model, and collapse it to an mDAG, representing its margin.

But the definition of the marginal model is implicit:

$$p(x_1, x_2, x_3, x_4) = \int p(u) p(x_1) p(x_2 | x_1, u) p(x_3 | x_2) p(x_4 | x_3, u) du$$

Actually determining whether or not a distribution satisfies the marginal Markov property **is hard**.

Model Description

We can write down a causal model, and collapse it to an mDAG, representing its margin.

But the definition of the marginal model is implicit:

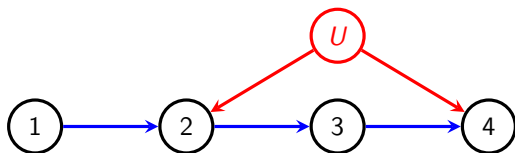
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Actually determining whether or not a distribution satisfies the marginal Markov property **is hard**.

Our strategy:

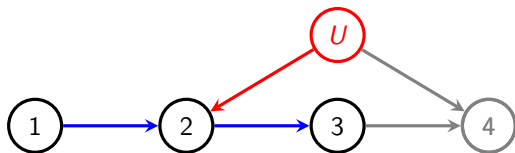
- derive some properties satisfied by the marginal model;
- define a new (larger) model that satisfies these properties;
- work with the larger model.

Ancestral Sets



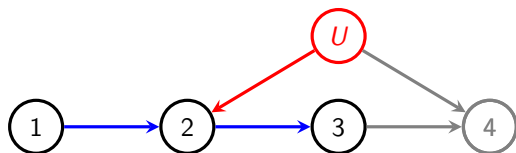
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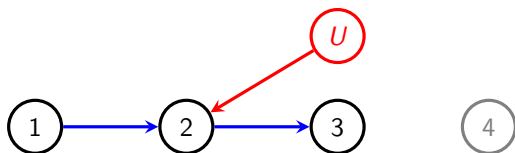
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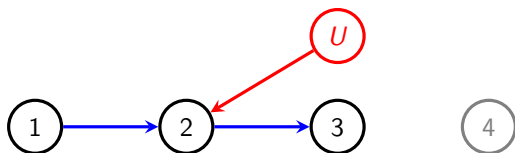
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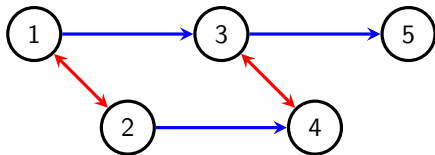


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Density has form corresponding to ancestral sub-graph.

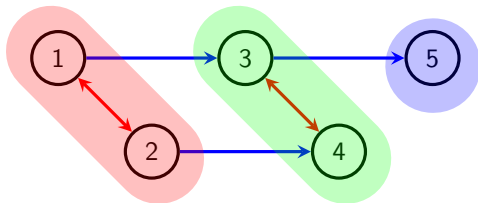
Factorization into Districts

District is a maximal set connected by latent variables / bidirected edges:



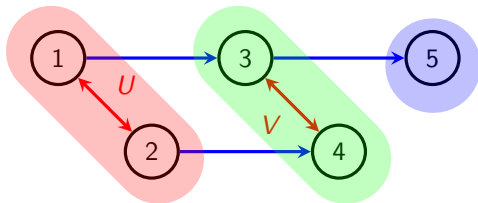
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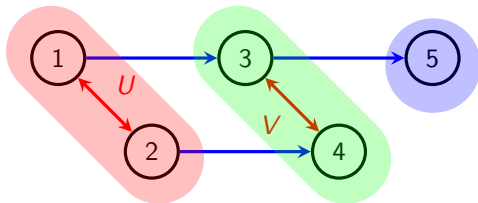
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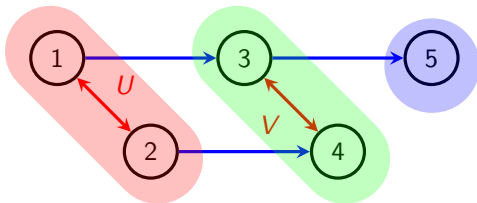
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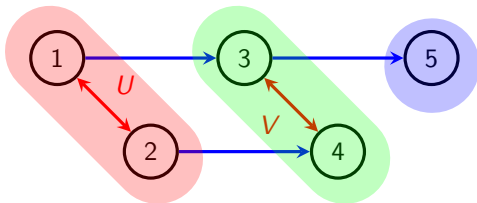
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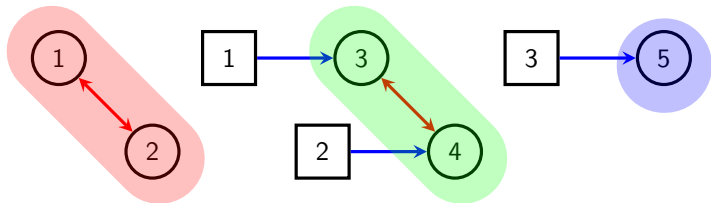
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 &= q_{12}(x_1, x_2) \cdot q_{34}(x_3, x_4 | x_1, x_2) \cdot q_5(x_5 | x_3) \\
 &= \prod_i q_{D_i}(x_{D_i} | x_{\text{pa}(D_i) \setminus D_i})
 \end{aligned}$$

Each q_D piece should come from the model based on district D and its parents ($\mathcal{G}[D]$).

Nested Model

We use these two rules to define our model.

Say (conditional) probability distribution p **recursively factorizes** according to mDAG \mathcal{G} and write $p \in \mathcal{N}(\mathcal{G})$ if:

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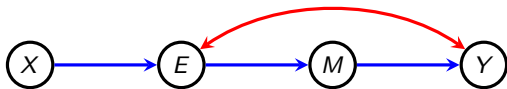
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Note that one can iterate between 1 and 2.

This defines the **nested Markov model** $\mathcal{N}(\mathcal{G})$. (Shpitser et al., 2014)

Example



Y childless,

Example



Y childless, so if $p \in \mathcal{N}(\mathcal{G})$, then

$$p(x, e, m) = p(x) \cdot p(e | x) \cdot p(m | e),$$

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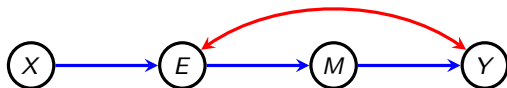


Y childless, so if $p \in \mathcal{N}(\mathcal{G})$, then

$$p(x, e, m) = p(x) \cdot p(e | x) \cdot p(m | e),$$

and therefore $X \perp\!\!\!\perp M | E$.

Example



Axiom 2:

$$p(x, e, m, y) = q_X(x) \cdot q_M(m | e) \cdot q_{EY}(e, y | x, m).$$

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This places a non-trivial constraint on p .

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We know

$$\mathcal{M}(\mathcal{G}) \subseteq \mathcal{N}(\mathcal{G}).$$

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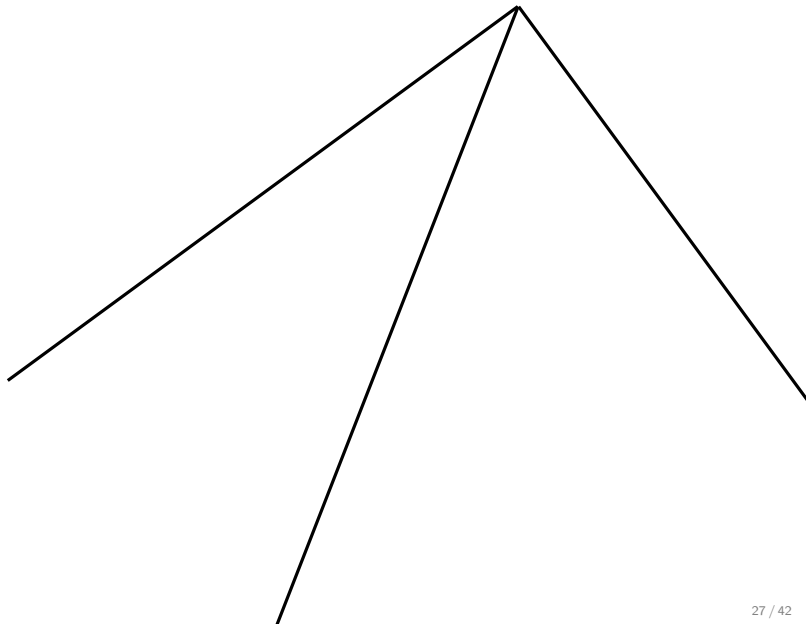
For discrete observed variables, the constraints implied by the nested Markov model are algebraically equivalent to causal model with latent variables (with suff. large latent state-space).

'Algebraically equivalent' = 'up to inequalities'.

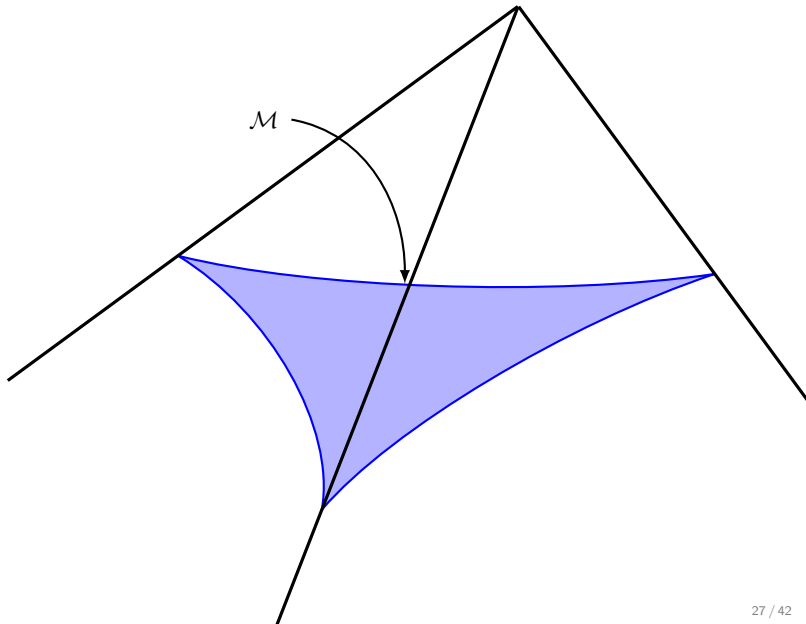
Any 'gap' $\mathcal{M}(\mathcal{G}) \subset \mathcal{N}(\mathcal{G})$ is due to inequality constraints.

So in particular they have the same dimension.

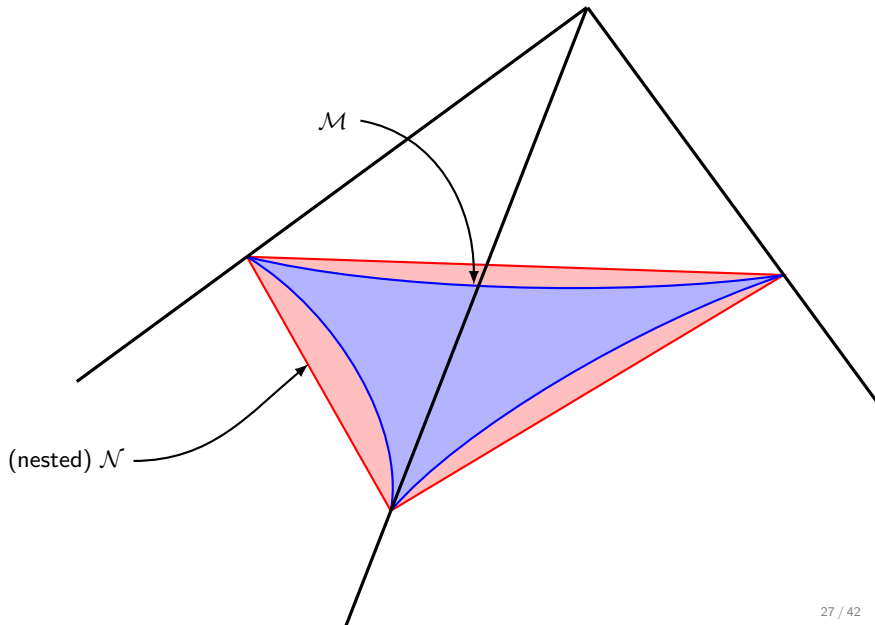
Getting the Picture



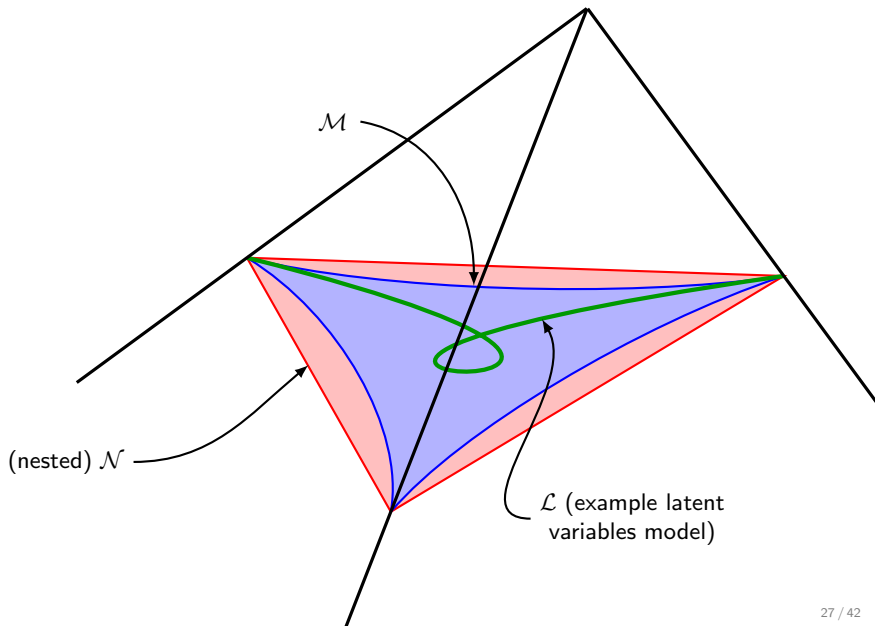
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Main Result

Nested model is a good approximation to the marginal model: in the discrete case it can be explicitly parameterized and fitted.

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Discrete nested models are curved exponential families.

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All parameters are of the form $p(\mathbf{X} \mid \text{do}(\mathbf{Y}))$: easily interpretable.

Wisconsin Data Example

Take only male respondents who were either drafted or didn't enter military at all (before 1975).

Continuous values dichotomised close to median.

Four binary indicators:

X family income $>$ \$5k in 1957;

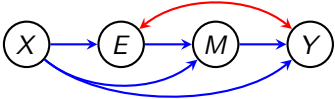
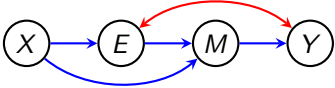
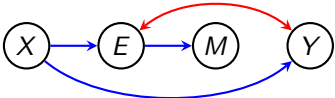
E education post high school;

M drafted into military;

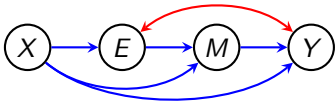
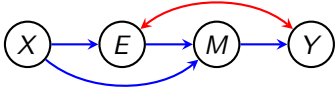
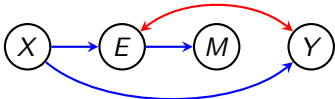
Y respondent income $>$ \$37k in 1992.

1,676 complete cases in 2^4 contingency table (minimum count 16).

Results

	model	deviance	d.f.
(a)		(saturated)	15
(b)		31.3	2
(c)		5.6	6

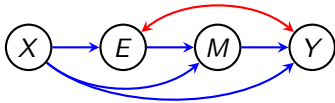
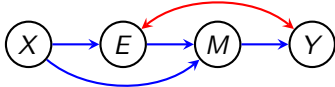
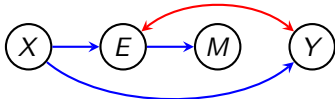
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Removing any edges from (c) strongly rejected.

Also find strong residual income effect:

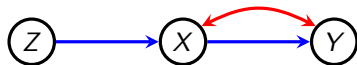
$$P(Y = 1 \mid \text{do}(X = 0)) = 0.36 \quad P(Y = 1 \mid \text{do}(X = 1)) = 0.50.$$

Outline

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- 2 An Example
- 3 mDAGs
- 4 Inequalities**
- 5 Testing, Fitting and Searching

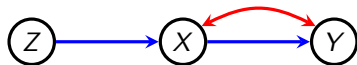
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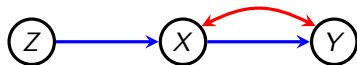


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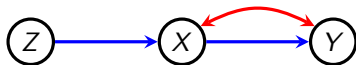
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Pearl (1995) showed that if the observed variables are discrete,

$$\max_x \sum_y \max_z P(X = x, Y = y | Z = z) \leq 1. \quad (*)$$

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e.g.

$$P(X = x, Y = 0 | Z = 0) + P(X = x, Y = 1 | Z = 1) \leq 1.$$

This is the **instrumental inequality**, and can be empirically tested.

Missing Edges Give Constraints

Proposition (Evans, 2012)

If X and Y are not joined by an edge in \mathcal{G} there is always a constraint induced on a discrete joint distribution.

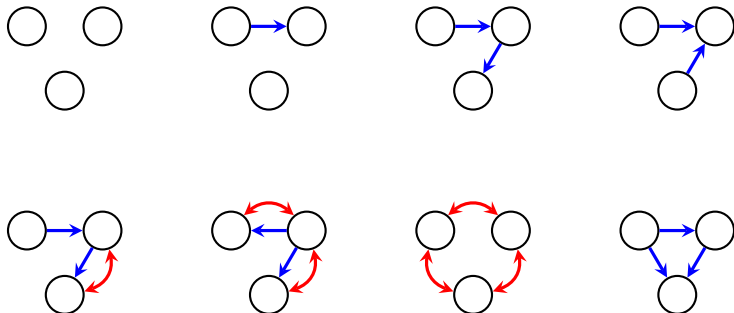
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Equivalence on Three Variables

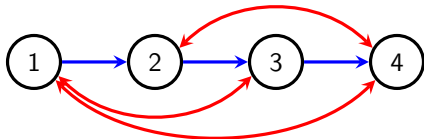
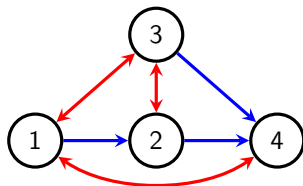
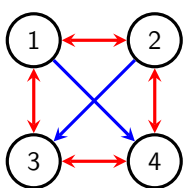
Markov equivalence (i.e. determining whether two models are observably the same) is hard.

Using Evans (2015) there are 8 unlabelled marginal models on three variables.



But Not on Four!

On four variables, it's still not clear whether or not the following models are saturated: (they are of full dimension in the discrete case)



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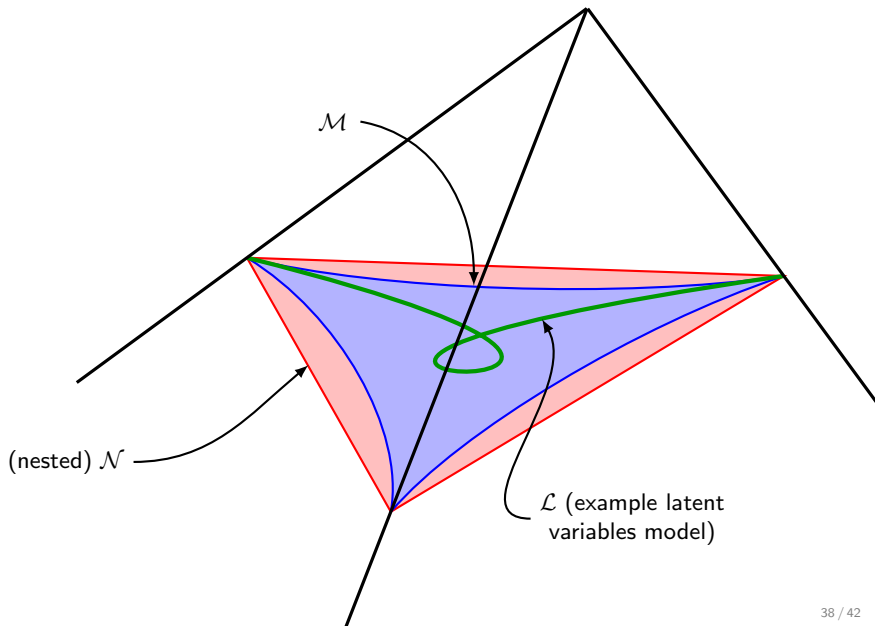
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- $\mathcal{L}(\mathcal{G}) \subseteq \mathcal{M}(\mathcal{G}) \subseteq \mathcal{N}(\mathcal{G})$.

So if we accept the latent variable model, or reject the nested model, same applies to marginal model.

That Picture Again



Some Extensions

We know nested models are curved exponential families, so justifies classical statistical theory:

- likelihood ratio tests have asymptotic χ^2 -distribution;
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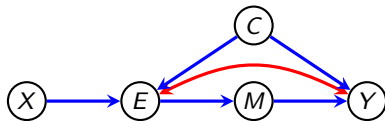
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Also, latent variable models **become** regular if state-space is large enough.

Can also include continuous covariates with outcome as multivariate response. e.g.:



Summary

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- Complete inequality constraints seem very complicated (though some hope exists);
- nice rule for model equivalence not yet available for either nested or marginal models.

Thank you!

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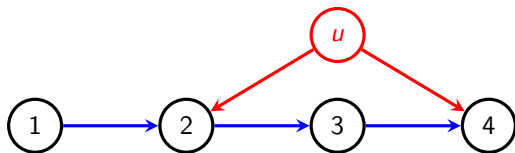
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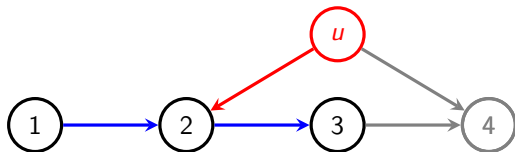
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Ancestral Sets



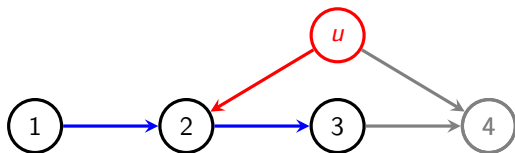
$$\begin{aligned} & p(x_1, x_2, x_3, x_4) \\ &= \sum_u p(u) p(x_1) p(x_2 | x_1, u) p(x_3 | x_2) p(x_4 | x_3, u) \end{aligned}$$

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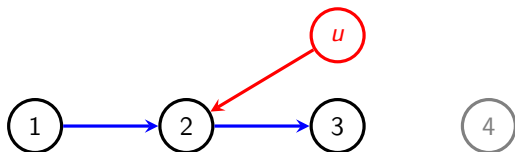
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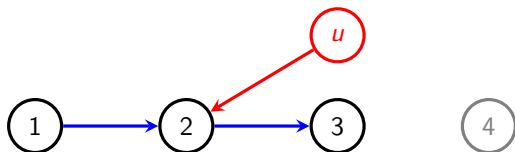
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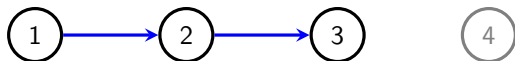
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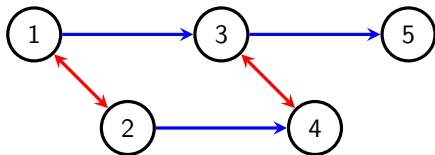


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Density has form corresponding to ancestral sub-graph.

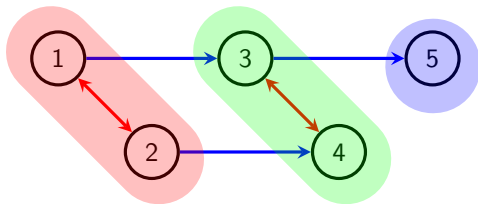
Factorization into Districts

District is a maximal set connected by latent variables / bidirected edges:



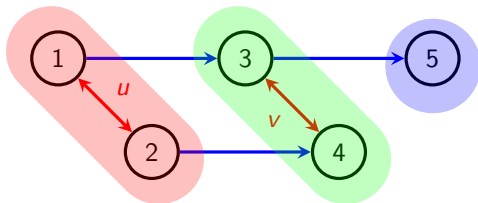
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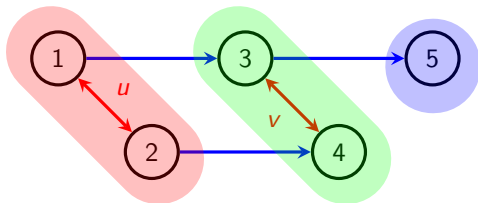
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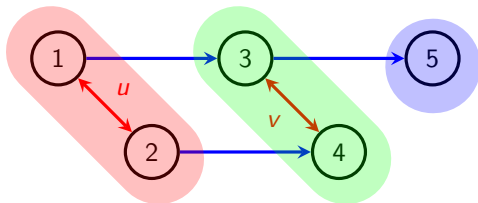
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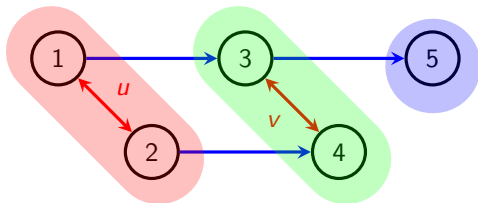
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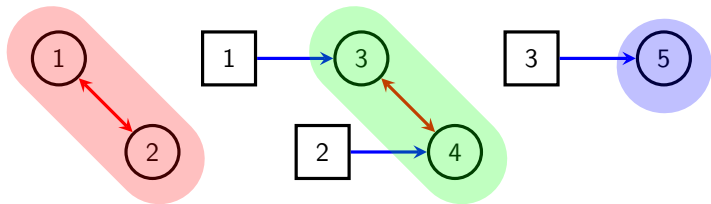
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Each q_D piece should come from the model based on district subgraph and its parents ($\mathcal{G}[D]$).

Nested Model

We use these two rules to define our model.

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Note that one can iterate between 1 and 2.

This defines the **nested Markov model** $\mathcal{N}(\mathcal{G})$.

Causal Coherence of mDAGs

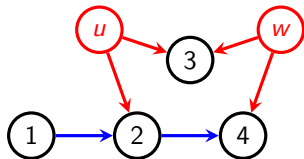
If P is represented by a DAG in a causally interpreted way, then intervening on some set of nodes $C \subseteq V$ can be represented by deleting incoming edges to C in \mathcal{G} . Call that graph $\mathcal{G}^{\bar{C}}$

Theorem (Evans, 2015)

If $C \subseteq O$ then $\mathbb{p}(\mathcal{G}^{\bar{C}}, O) = \mathbb{p}(\mathcal{G}, O)^{\bar{C}}$; i.e. the projection respects causal interventions.

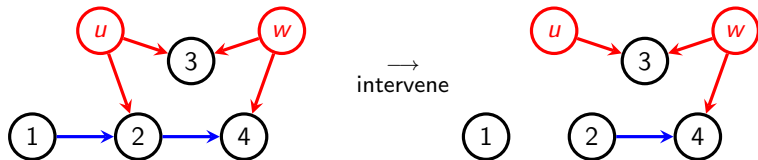
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If we intervene on some observed variables, this 'breaks' their dependence upon their parents.



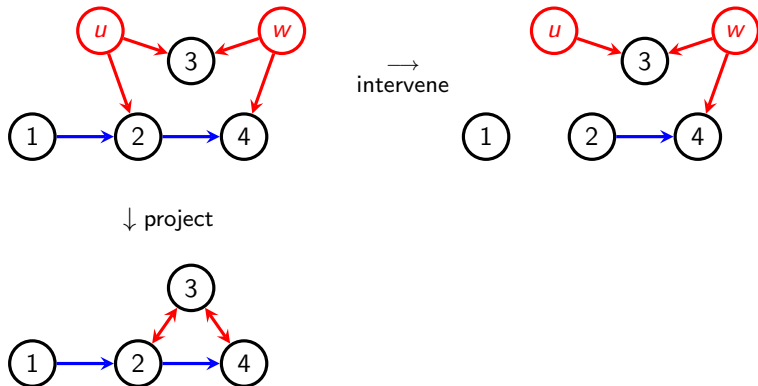
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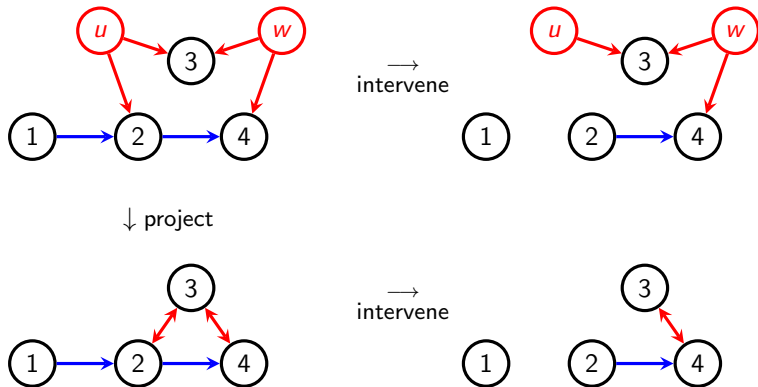
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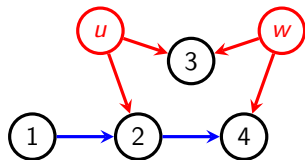
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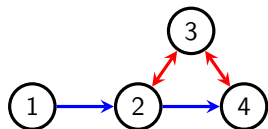


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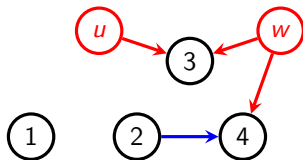
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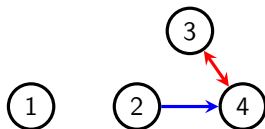
↓ project



→
intervene

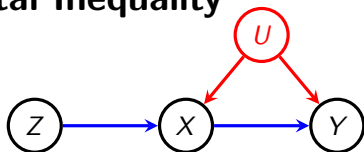


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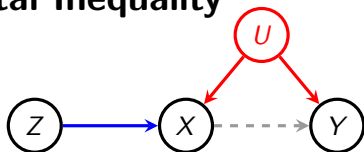
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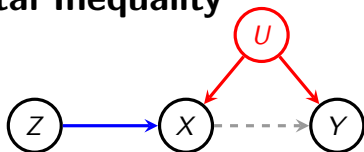
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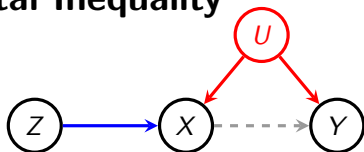
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Can't observe p^* but:

- **Consistency:** $p(\xi, y | z) = p^*(\xi, y | z)$ for each z, y ; and
- **Independence:** $Y \perp\!\!\!\perp Z$ under p^* .

Solution: A Different Proof

For each $x = \xi$ we require p_ξ^* :

$$p_\xi(\xi, y | z) = p_\xi^*(\xi, y | z) \text{ for each } y, z, \quad Y \perp\!\!\!\perp Z [p_\xi^*].$$

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We say that the probabilities $p(x, y | z)$ are **compatible** with $Y \perp\!\!\!\perp Z$.

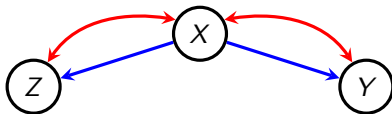
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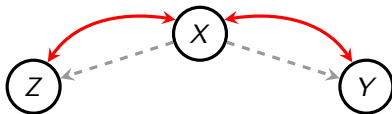
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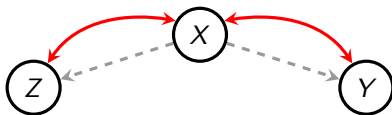


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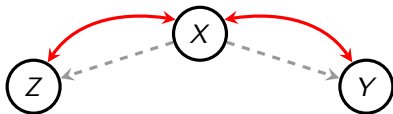
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So by the same argument, for fixed ξ , $p(\xi, y, z)$ must be compatible with a (fictitious) distribution p_{ξ}^* in which $Y \perp\!\!\!\perp Z$.

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How does this help us with other graphs?

The argument works precisely because cutting edges led to an independence:



Z is independent of Y in the graph after cutting edges emanating from X .

So by the same argument, for fixed ξ , $p(\xi, y, z)$ must be compatible with a (fictitious) distribution p_{ξ}^* in which $Y \perp\!\!\!\perp Z$.

[Note for the IV model, the conditional distribution $p(\xi, y | z)$ had to be compatible.]

d-Separation

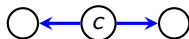
A **path** is a sequence of edges in the graph; vertices may not be repeated.

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A path from v to w is **blocked** by $C \subseteq V \setminus \{v, w\}$ if either

(i) any non-collider is in C :

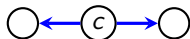


d-Separation

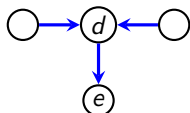
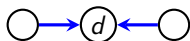
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(ii) or any collider is not in C , nor has descendants in C :

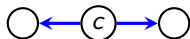


d-Separation

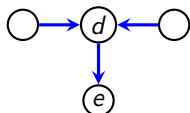
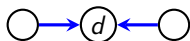
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Two vertices v and w are **d-separated** given $C \subseteq V \setminus \{v, w\}$ if **all** paths are blocked.