

Graphical Models and the PC Algorithm

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The Problem with Causality

“Causality is the centerpiece of the universe”¹

“The central aim of many studies . . . is the elucidation of cause-effect relationships between variables or events”²

- Criticism of statistical science: focus on probabilistic and statistical inference at the expense of causal enquiry

¹Causality - Wikipedia, the free encyclopedia

²Preface to *Pearl (2000)*

Conditional Independence

Definition (Conditional Independence)

The random variables X and Y are said to be conditionally independent given the value of a third random variable Z , if $f(X|Y, Z) = f(X|Z)$.

- Write: $X \perp\!\!\!\perp Y \mid Z$
- Intuitively, if Z is known, Y adds no information about the value of X .
- The difference between independence and conditional independence is demonstrated by the Yule-Simpson Paradox.

Example: The Berkeley sex-bias case

The University of California, Berkeley, were sued for bias against women applying to grad school:

- In the university as a whole, men were more likely to be admitted to a course than women
- Examining individual departments (conditioning on the departments), there was no significant bias against women—in fact, most departments showed a slight bias against men
Explanation:
 - women tended to apply for courses with low admission rates
 - men tended to apply for courses with high admission rates

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Yule-Simpson Paradox

Let n_{ij}, N_{ij} , $i \in \{1, 2\}$ and $j \in \{A, B\}$, be integers. Then it is possible that:

$$\frac{n_{1A}}{N_{1A}} < \frac{n_{1B}}{N_{1B}}$$

and

$$\frac{n_{2A}}{N_{2A}} < \frac{n_{2B}}{N_{2B}}$$

but

$$\frac{n_{1A} + n_{2A}}{N_{1A} + N_{2A}} > \frac{n_{1B} + n_{2B}}{N_{1B} + N_{2B}}$$

Applying this to the calculation of conditional probabilities leads to the Yule-Simpson paradox, credited to George Udny Yule (1903) and popularised by E.H. Simpson (1951).

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- Nodes:** The vertices ($i \in V$) of the graph
(Nodes and vertices used interchangeably)
- Edges:** Connections ($(i, j) \in E$) between vertices
- Path:** A route along (directed) edges from one node to another
(e.g. $i \rightarrow j \rightarrow k \rightarrow l$)

Definition (Graphical Model)

A graphical model G is a system of nodes and connecting edges:
 $G = (V, E)$

The role of graphs in probabilistic and statistical modeling is threefold:

- 1 to provide convenient means of expressing substantive assumptions;
- 2 to facilitate economical representation of joint probability functions; and
- 3 to facilitate efficient inferences from observations.

Conditional Independence Graph

Definition (Conditional Independence Graph)

The conditional independence graph of X is the **undirected** graph $G = (V, E)$ where $V = \{1, 2, \dots, n\}$ and (i, j) is not in the edge set E iff $X_i \perp\!\!\!\perp X_j \mid X_{V \setminus \{i, j\}}$.

More informally:

- Start with the complete graph, where each node is connected to all other nodes
- Remove the edge between X_i and X_j if

$$X_i \perp\!\!\!\perp X_j \mid \text{rest}$$

N.B.: The conditional dependencies do not represent causal or directed relationships between variables.

The Pairwise Markov Property

A graph has the pairwise Markov property if, for all non-adjacent (not directly connected) vertices i and j ,

$$X_i \perp\!\!\!\perp X_j \mid X_{V \setminus \{i, j\}}$$

- Undirected conditional independence graphs are formed using this definition

Therefore, if X_i and X_j are non-adjacent vertices:

- they are independent conditional on the remaining nodes
- X_j is irrelevant for the prediction of X_i , and vice-versa
- *Separation Theorem:* $X_i \perp\!\!\!\perp X_j \mid \text{rest} \Rightarrow X_i \perp\!\!\!\perp X_j \mid X_a$, where X_a are the vertices separating X_i and X_j .

The Local Markov Property

A graph has the local Markov property if, for every vertex i , with boundary $a = \text{bd}(i)$ and b the set of remaining vertices,

$$X_i \perp\!\!\!\perp X_b \mid X_a$$

More informally, if:

$$X_i \perp\!\!\!\perp \text{rest} \mid \text{boundary}$$

- Closely related to prediction—conditioned only on adjacent variables

The Global Markov Property

Let a , b and c be disjoint subsets of V . Then, a graph has the global Markov property if, whenever b and c are separated by a in the graph, then:

$$X_b \perp\!\!\!\perp X_c \mid X_a$$

- Global in the sense that the subsets are potentially arbitrary

Equivalence of Markov Properties

The three Markov properties: pairwise Markov, local Markov and global Markov, are equivalent.

- As the boundary set is always a separating set, global Markov \Rightarrow local Markov
- Local Markov \Rightarrow pairwise Markov
- By separation theorem, pairwise Markov \Rightarrow global Markov

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Definition (Directed Acyclic Graph)

A graph $G = (V, E)$ is called a directed acyclic graph if all edges are directed and there are no cycles (i.e. it is impossible to return to any point).

- $X \rightarrow Y \implies X$ "causes" Y
- Various theorems—and background information—can be used to identify which conditional dependencies are causal in nature.
- Independent variables (i.e. no directed edge) may be dependent conditional on the remaining variables (Berkson's Paradox)

1 Serial Connection

A series of nodes: $i \rightarrow j \rightarrow k$

2 Diverging Connection

One node leads to several: $j \leftarrow i \rightarrow k$

3 Converging Connection

Several nodes lead to one path: $j \rightarrow i \leftarrow k$

Definition (d -separation)

A set Z is said to d -separate (directionally separate) X from Y iff Z blocks every path from a node in X to a node in Y

Properties of a DAG

Definition (Faithfulness)

A distribution P is faithful to a DAG D if the all conditional independence relations for P can be derived from d -separation.

- Faithful graphs can be estimated using conditional independence relations
- Direction means that the graph is conditioned only on previous nodes
- Directed independence graphs are therefore based on the local and not pairwise Markov property

Definition (Skeleton of a DAG)

The graph generated by replacing all directed edges of a DAG with undirected edges is called a skeleton.

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Estimating DAG Structures

Suppose we have a multivariate data sample and assume:

- p variables and sample size n
- $X \sim N_p(\mu, \Sigma)$
- This multivariate normal distribution is faithful
- The underlying graph is sparse (i.e. not too many edges)

Then, the structure of a DAG can be recovered using conditional independence relations.

Pairwise vs. Local Markov Property

Estimate the skeleton using the pairwise Markov, not the local Markov property:

- For any given vertex, there are 2^{p-1} ways of partitioning the remaining vertices into "boundary" and "rest" groups
- If p is large (or $p > n$), this is both computationally and statistically infeasible
- In contrast, the pairwise property has only $(k - 1)$ ways of partitioning the remaining vertices

Conditional Independence

Definition (Partial Correlation)

For $i \neq j \in 1, \dots, p$, $k \in \text{rest}$, let $\rho_{i,j|k}$ be the partial correlation between X_i and X_j given X_r ; $r \in k$.

- As the distribution is multivariate normal, $X_i \perp\!\!\!\perp X_j \mid X_r \iff \rho_{i,j|k} = 0$
- A test for conditional independence is therefore a test for partial correlation between the variables
- The partial correlations can be estimated, for example, via regression

Test for Conditional Independence

Definition (Fisher's Z-Transform)

Let:

$$Z(i, j|k) = \frac{1}{2} \left(\frac{1 + \hat{\rho}_{i,j|k}}{1 - \hat{\rho}_{i,j|k}} \right)$$

Then:

$$\sqrt{n - |k| - 3} |Z(i, j|k)| \sim N(0, 1)$$

- Test for independence using classical test at significance level α
- Kalisch and Bühlmann show that the choice of α is not too important
- Various other tests are available, using different approaches and for different distributions

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Start with the complete undirected graph, \tilde{C} with vertices $V = X_1, \dots, X_p$. Then:

- 1 Set $\ell = -1$ and $C = \tilde{C}$
- 2 Increase ℓ by one. For all pairs of adjacent nodes:
 - ▶ Check for conditional independence
 - ▶ Remove edge (X_i, X_j) if $X_i \perp\!\!\!\perp X_j \mid \text{rest}$
- 3 Repeat step 2 until $\ell = m$ or until each node has fewer than $\ell - 1$ neighbours

Stopping level m

Let $m, each \in \max \ell, m$ denote the stopping level of the algorithm and q be the maximum number of neighbours. It can be shown that:

- 1 The PC Algorithm constructs the true skeleton of the DAG
- 2 The stopping level is $m, each \in q - 1, 1$

Consistency of the PC Algorithm I

Let G be a DAG with probability distribution P . The following assumptions are made:

- 1 The distribution is multivariate normal and is faithful w.r.t. G
- 2 The dimension is $p_n = O(n^a)$, $a \geq \infty$
→ high dimensionality
- 3 The maximum number of neighbours, $q_n = O(n^{1-b})$, $0 < b \leq 1$
→ the graph is sparse
- 4 The partial correlations (absolute values) are bounded from above and below:
→ a regularity condition

Consistency of the PC Algorithm II

Denote by G_{skel} the true skeleton of a DAG G , and let the estimate from the PC Algorithm be \hat{G}_{skel} . Then, under the above assumptions, it can be shown that, for some $C \geq 0$:

$$P(\hat{G}_{skel} = G_{skel}) = 1 - O(\exp(-Cn^{1-2d})) \rightarrow 1, n \rightarrow \infty$$

Additionally, the stopping level is data dependent,

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Example using Simulated Data

Construct an adjacency matrix describing the conditional independence relations contained in a randomly generated graph of dimension p .

- Begin with a matrix of zeroes (i.e. no edges)
- Independent realisations of a Bernoulli random variable with parameter s determine which edges are connected. Call s the sparseness of the model
- For the edges in the graph (ones in the adjacency matrix), independent realisations of a Uniform[0.1, 1] distribution are used to model the partial correlations

Then, $X_1 = \epsilon_1 \sim N(0, 1)$, and the remaining nodes are calculated recursively as follows:

$$X_i = \sum_{k=1}^{i-1} A_{ik} X_k + \epsilon_i \quad i = 2, \dots, p$$

Summary Statistics

The PC algorithm is to be compared with two alternative methods:

- Greedy Equivalent Search (GES)
- Maximum Weight Spanning Trees (MWST)

The following statistics allow their characteristics to be compared:

- TDR** True discovery rate, the proportion of edges in the estimated model that are edges in the true model
- FPR** False positive rate, the proportion of edges in the estimated model that have been falsely identified
- TPR** True positive rate, the proportion of true edges that have been identified by the model

Method	ave[TPR]	ave[FPR]	ave[TDR]
PC	0.57 (0.06)	0.02 (0.01)	0.91 (0.05)
GES	0.85 (0.05)	0.13 (0.04)	0.71 (0.07)
MWST	0.66 (0.07)	0.06 (0.01)	0.78 (0.06)

The PC algorithm:

- achieves much higher TDR than GES or MWST
- identifies a lower proportion of the true nodes, but also has fewer false positives

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