

The Query Complexity of Verifying a Causal Graph from Single-Node Interventions

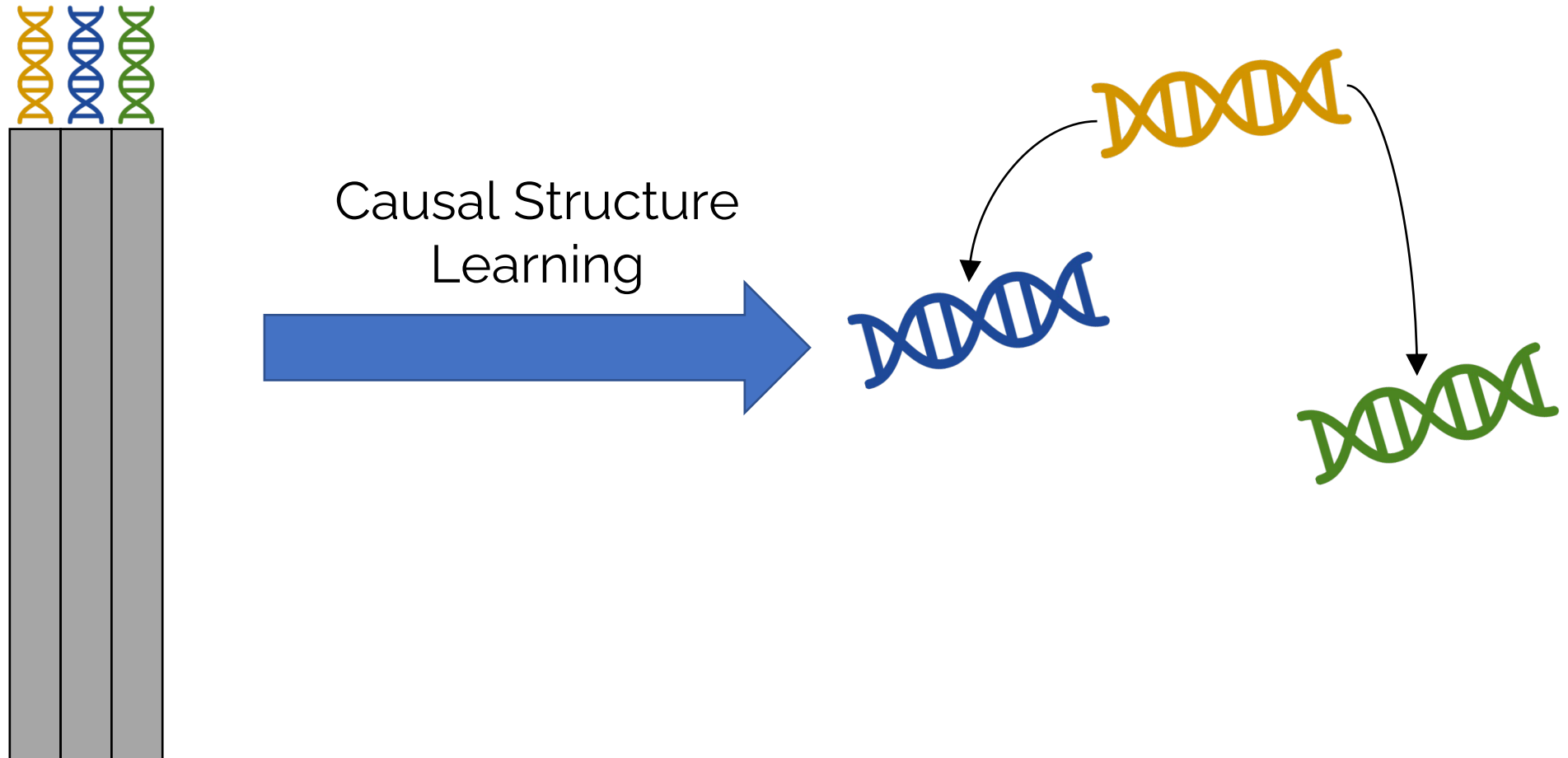
or: How hard is it to learn a causal model?

Chandler Squires

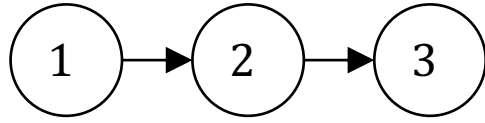
Causal Structure Learning (aka, Causal Discovery)

An inverse problem: given data, find the causal graph

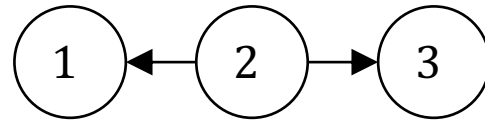
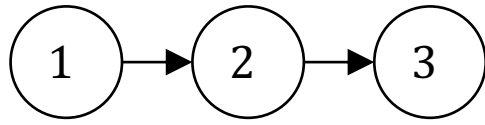
A Canonical Application: Learning Gene Regulatory Networks



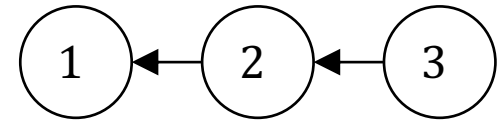
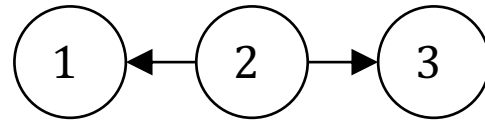
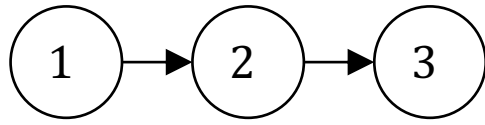
Observational data is (in general) insufficient
to identify the true graph



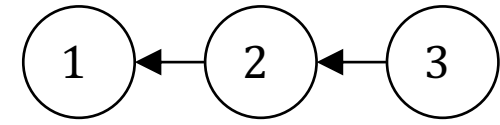
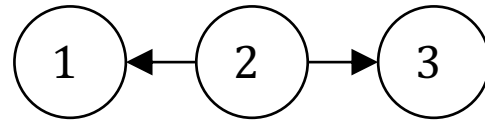
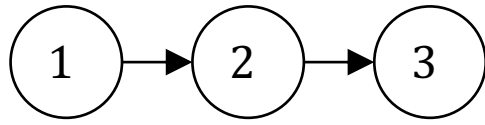
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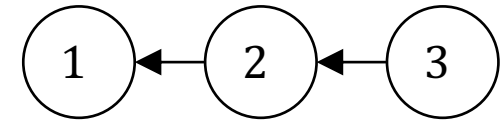
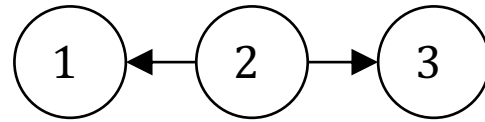
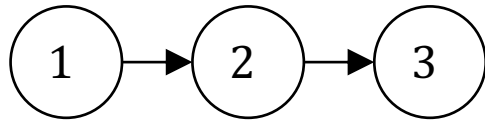
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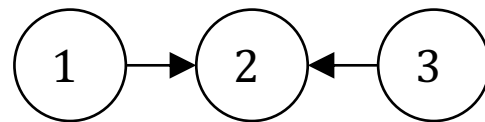
the **only**¹ Level 1 property (i.e., at the level of $\mathbb{P}(X_1, X_2, X_3)$) entailed by these graphs is $X_1 \perp\!\!\!\perp X_3 \mid X_2$.

¹without further parametric assumptions

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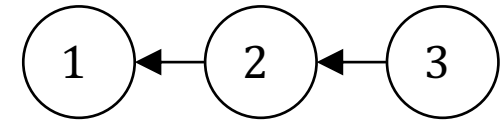
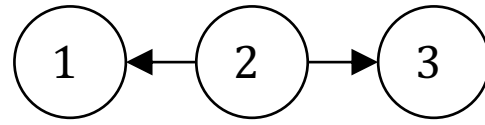
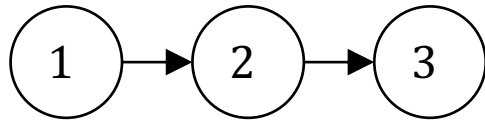
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However, this graph implies that $X_1 \perp\!\!\!\perp X_3$

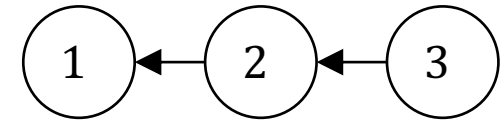
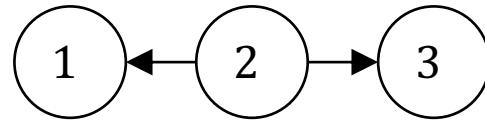
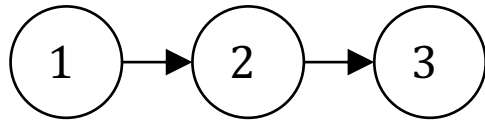
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What can we identify from observational data?



We call graphs which imply the same set of conditional independences **Markov equivalent**.

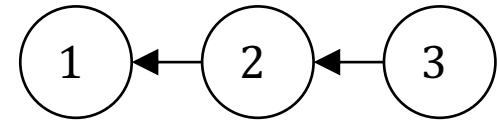
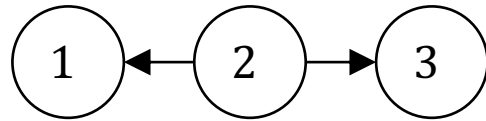
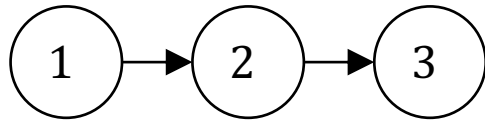
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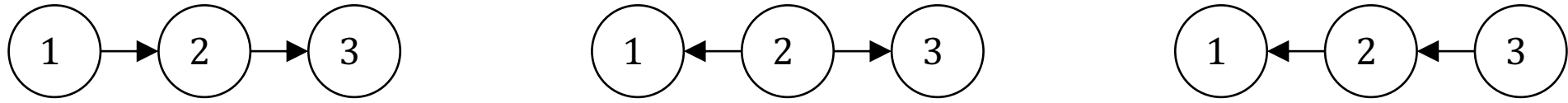
From observational data, we can only identify up to the **Markov equivalence class**.

What can we identify from observational data?

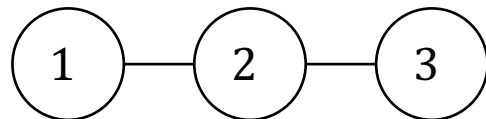


Two graphs are Markov equivalent if and only if they have the same adjacencies and **v-structures** (patterns of the form $i \rightarrow k \leftarrow j$ with i and j non-adjacent).

What can we identify from observational data?

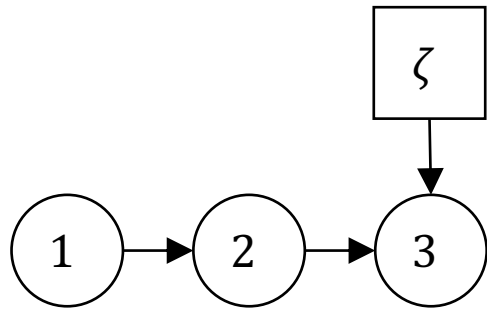


- We represent the Markov equivalence class by the **essential graph**, which:
- Has the same adjacencies as the members of the Markov equivalence class
 - Only has an edge directed if it is directed the same way across the entire Markov equivalence class.



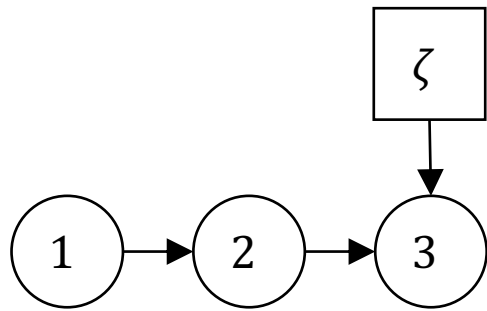
What do we learn from
interventions?

Single-node interventions teach us the directions of edges adjacent to the intervened node

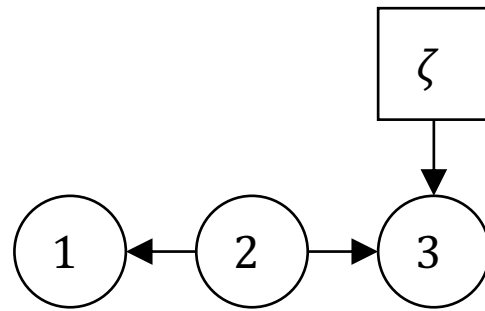


Only X_3 is affected

Single-node interventions teach us the directions of edges adjacent to the intervened node

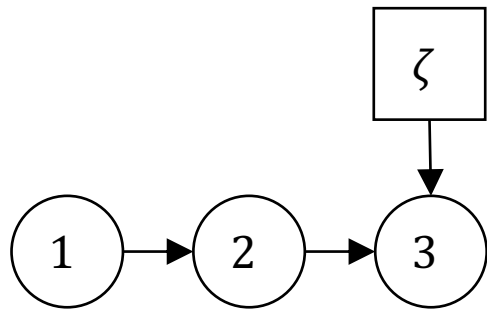


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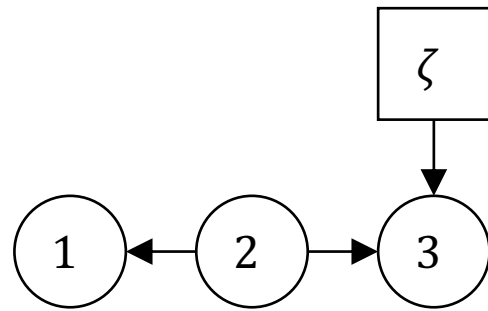


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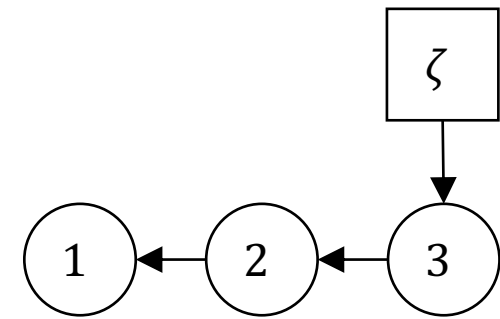
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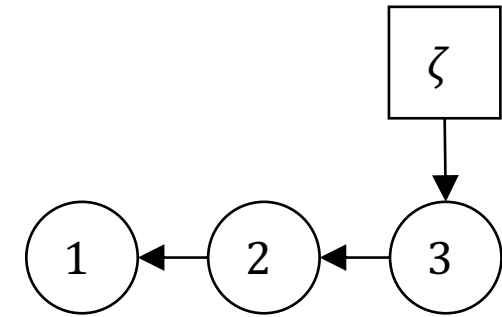
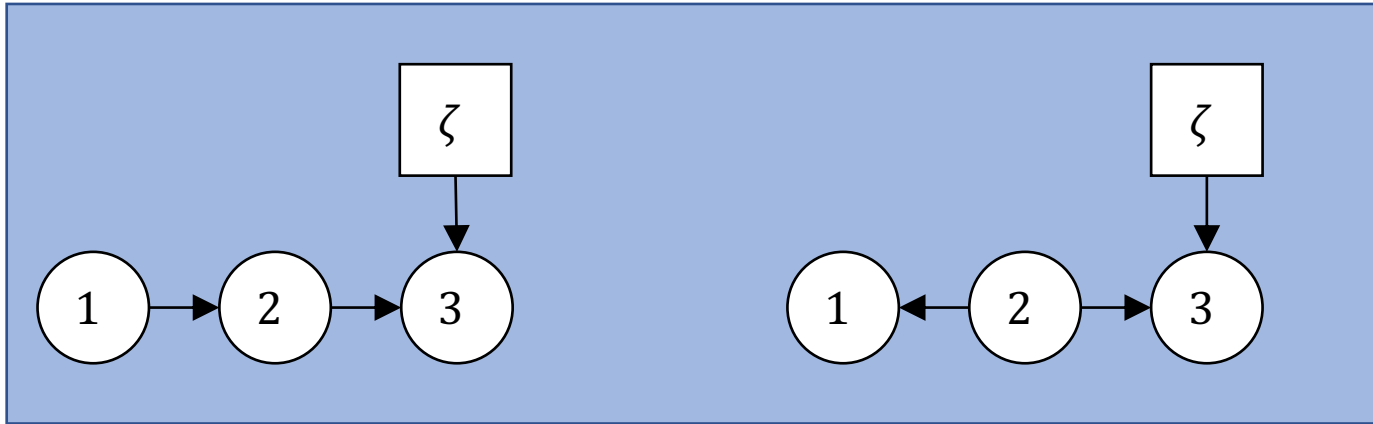


Only X_3 is affected

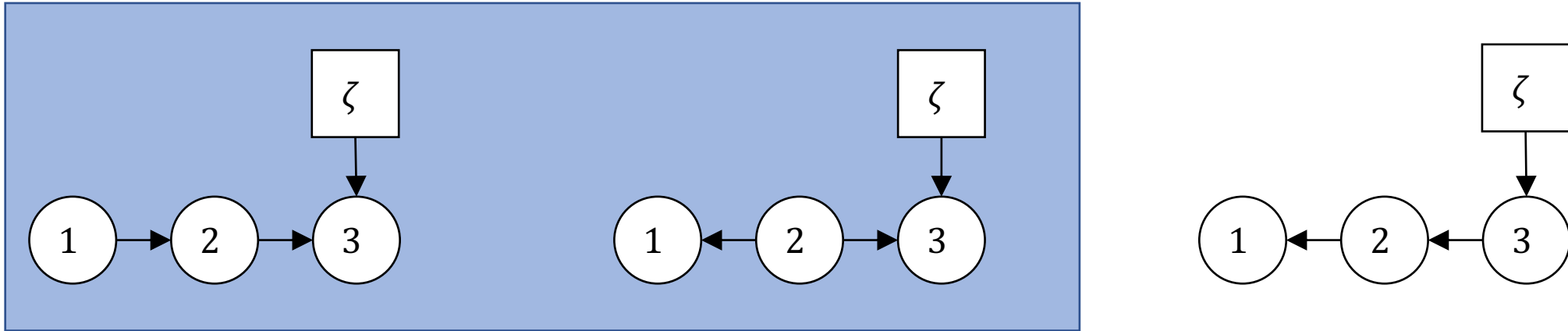


All three variables are affected

Interventional Markov equivalence



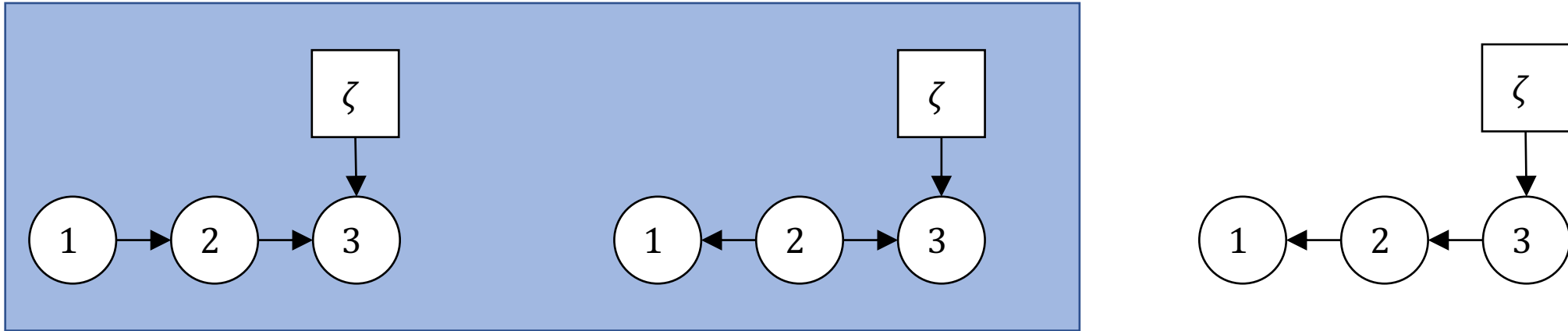
Interventional Markov equivalence



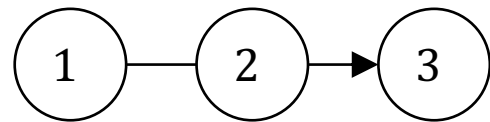
Denote the set of interventions as $\mathcal{J} = \{\{3\}\}$

We refer to the set of graphs entailing the same conditional independences and **conditional invariances** under \mathcal{J} as the **\mathcal{J} -Markov equivalence class**.

Interventional Markov equivalence

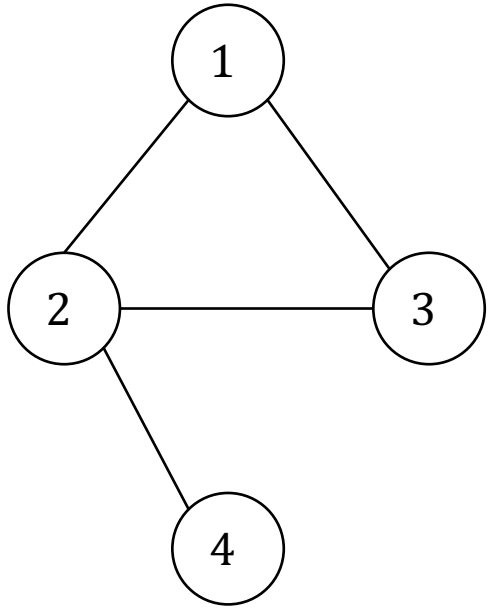


We represent the \mathcal{J} -Markov equivalence class using the **\mathcal{J} -essential graph**.

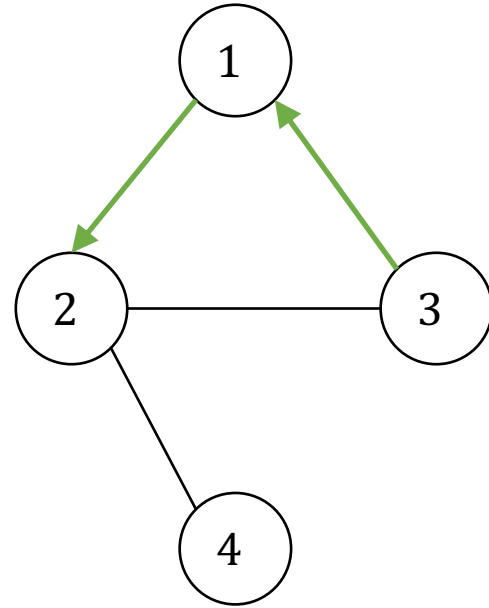
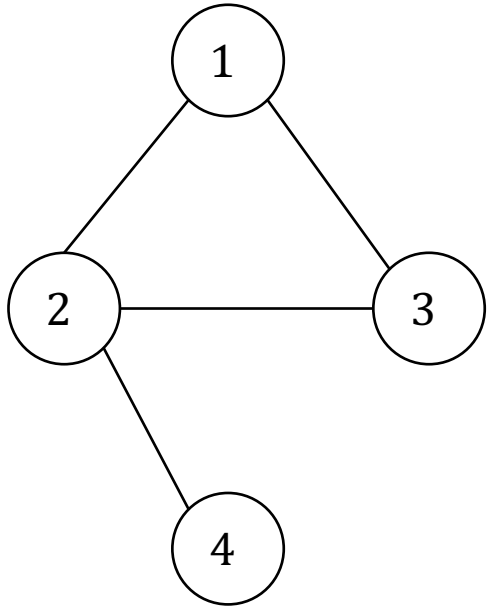


Do interventions teach us any
other orientations?

Yes: through the application of Meek's orientation rules

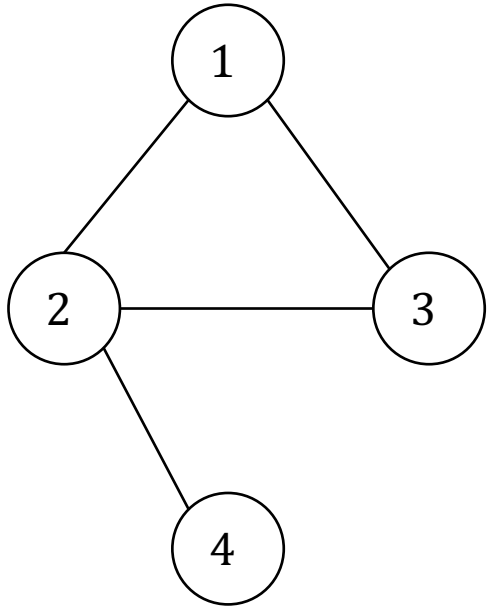


From observational data

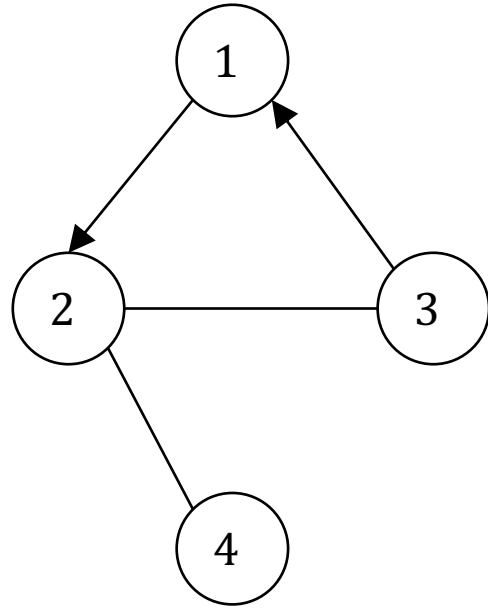


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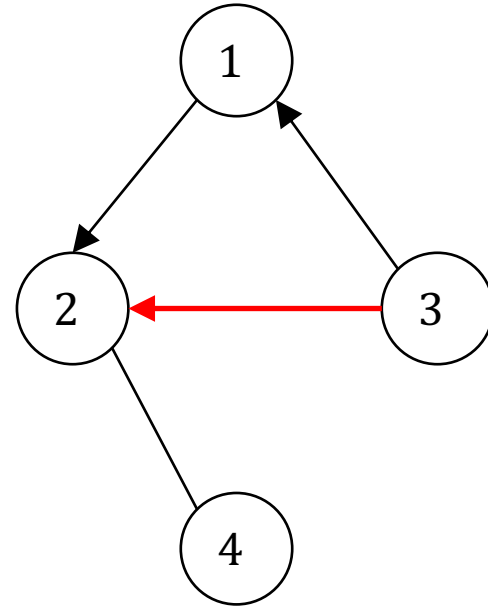
Adjacencies from an intervention on X_1



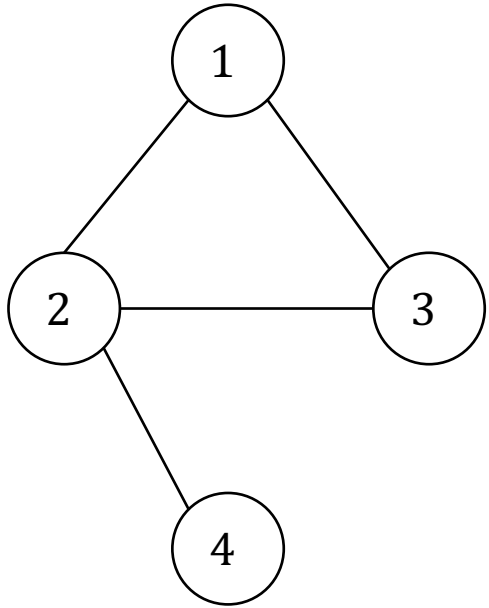
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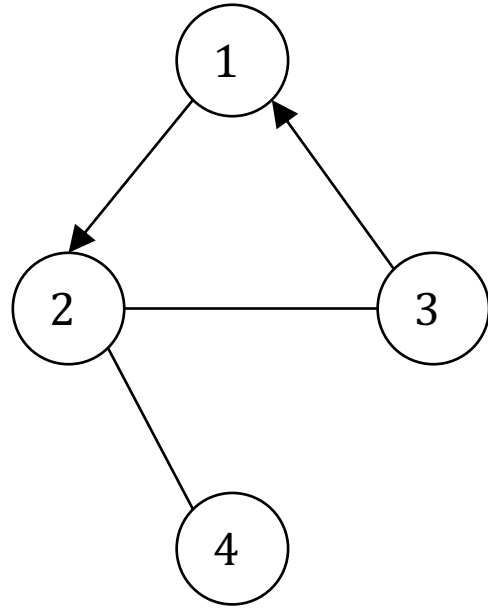
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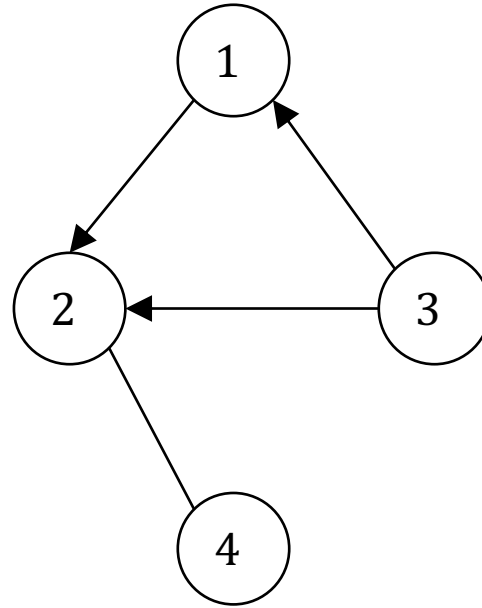
Avoid cycles
(Meek rule 2)



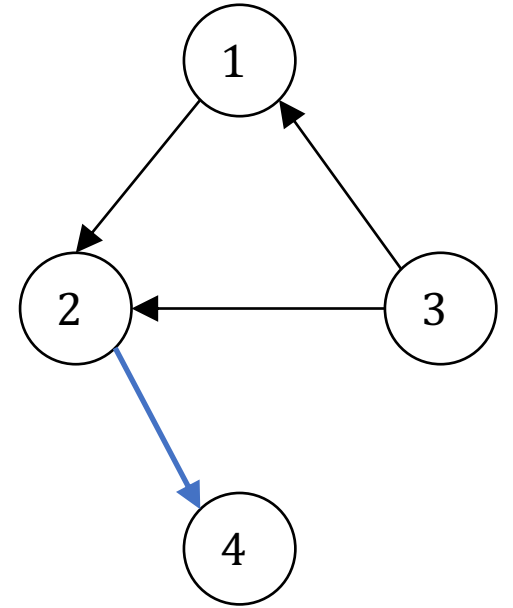
From observational data



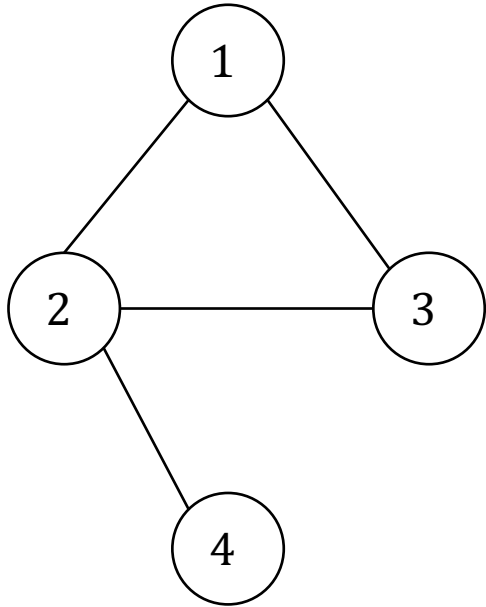
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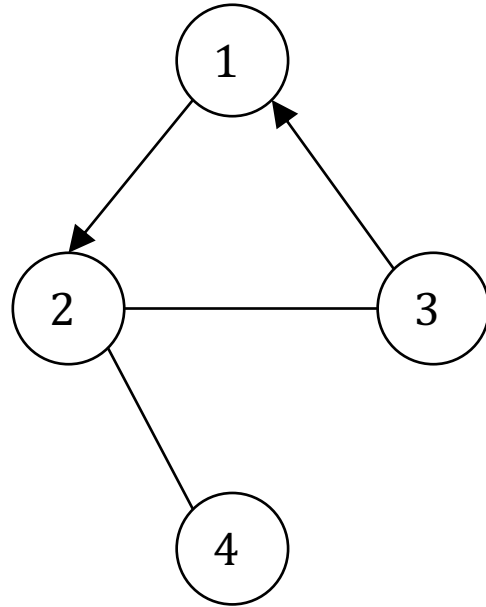
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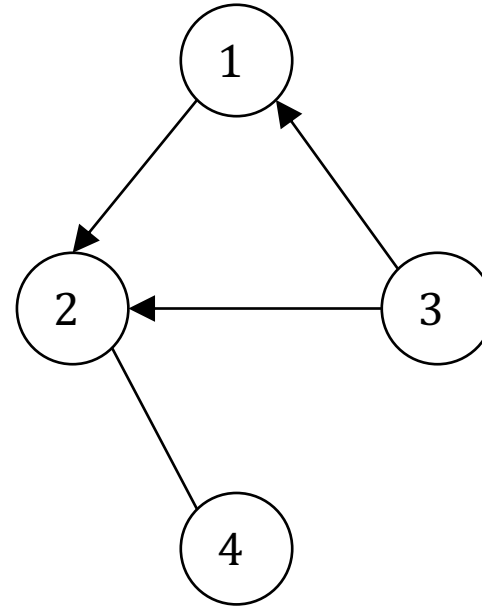
Avoid new v-structures
(Meek rule 1)



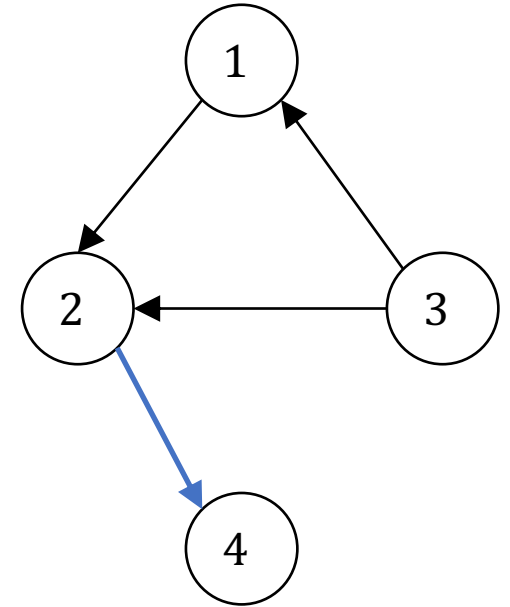
From observational data



Adjacencies from an intervention on X_1



Avoid cycles
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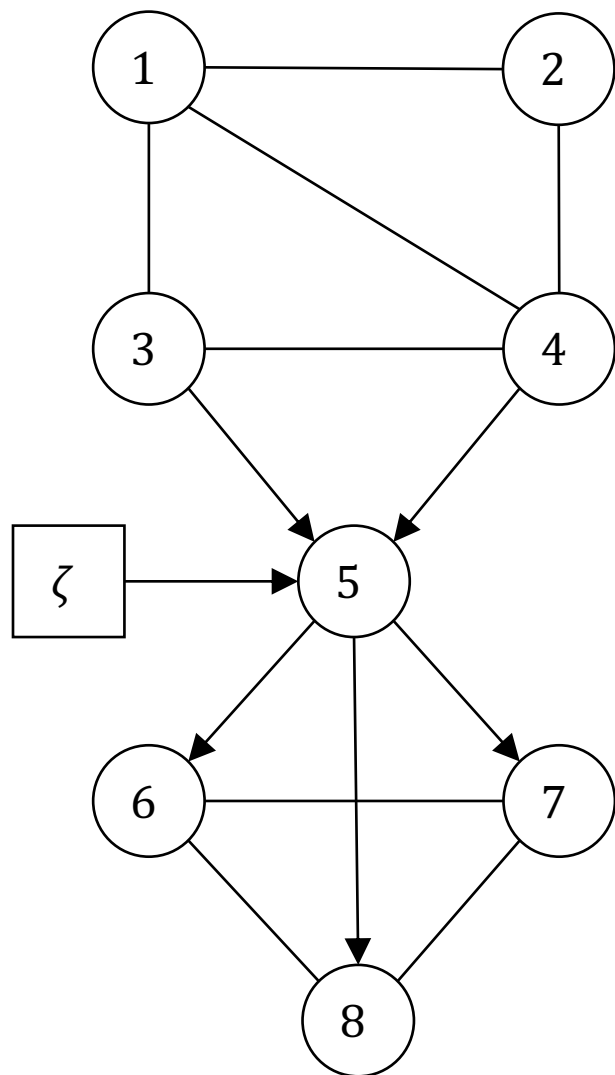


Avoid new v-structures
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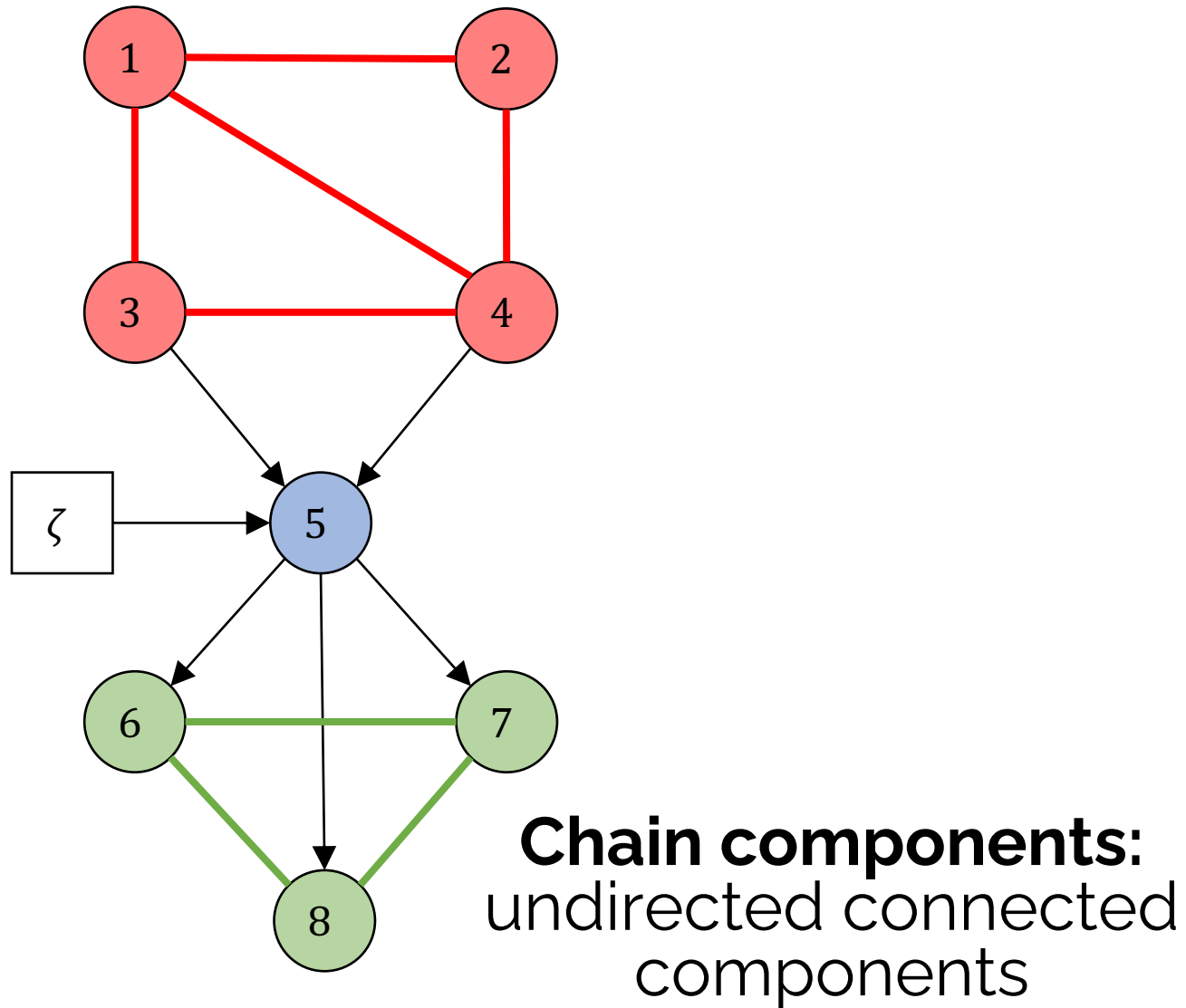
Note: There are two other Meek orientation rules; together these four rules are complete.

How many single-node interventions are required to fully orient a DAG?

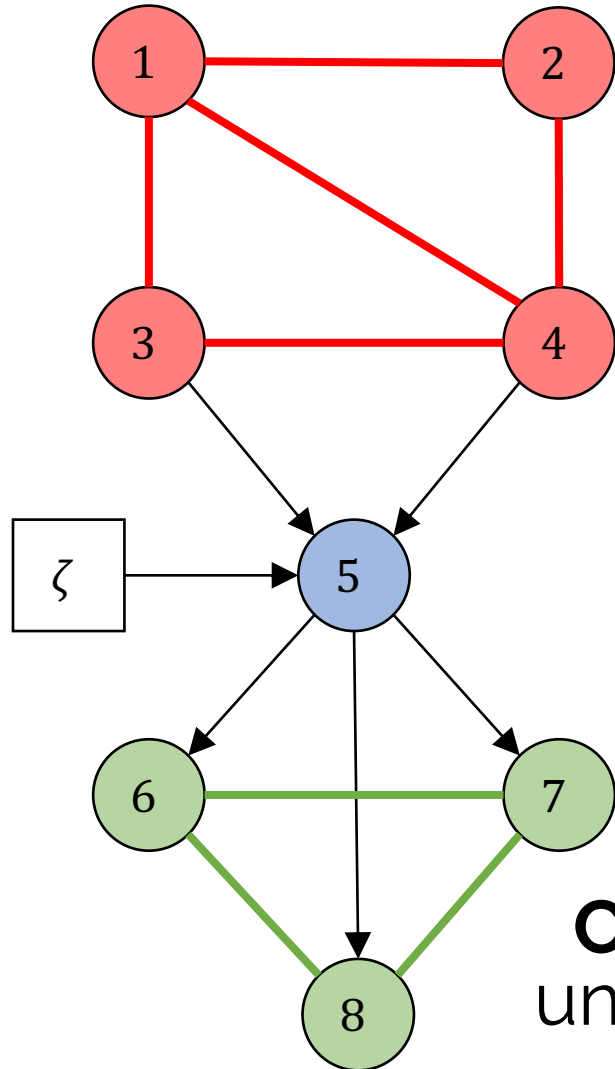
\mathcal{J} -essential graphs decompose nicely



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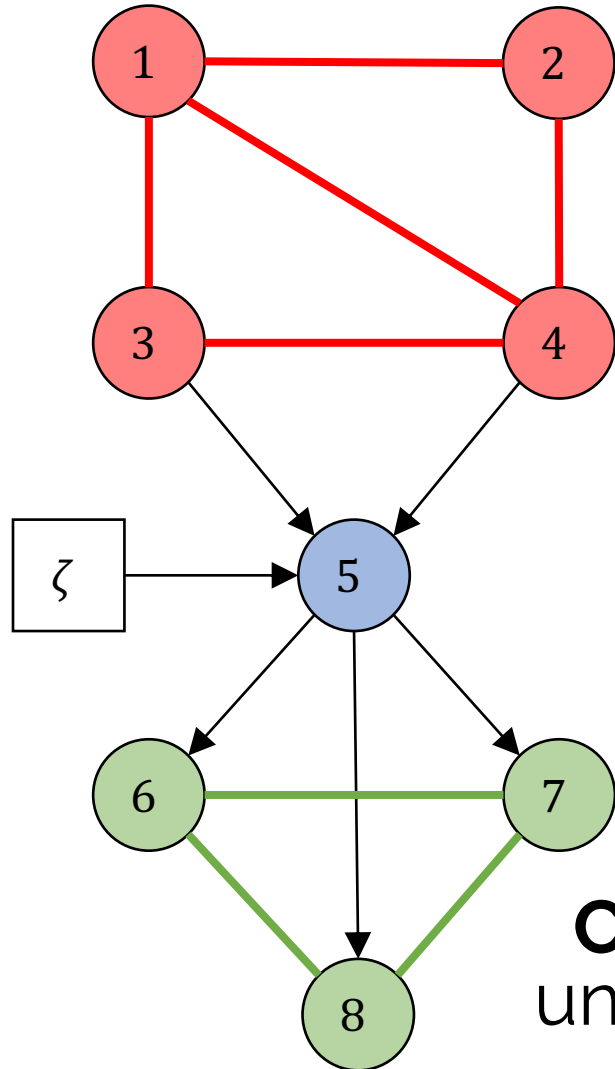
\mathcal{J} -essential graphs decompose nicely



- The chain components are **chordal**.
 - i.e., there are no cycles of length ≥ 4 which don't have a chord.

Chain components:
undirected connected
components

\mathcal{J} -essential graphs decompose nicely



- The chain components are **chordal**.
 - i.e., there are no cycles of length ≥ 4 which don't have a chord.
- The orientations within separate chain components are **independent**.

Chain components:
undirected connected
components

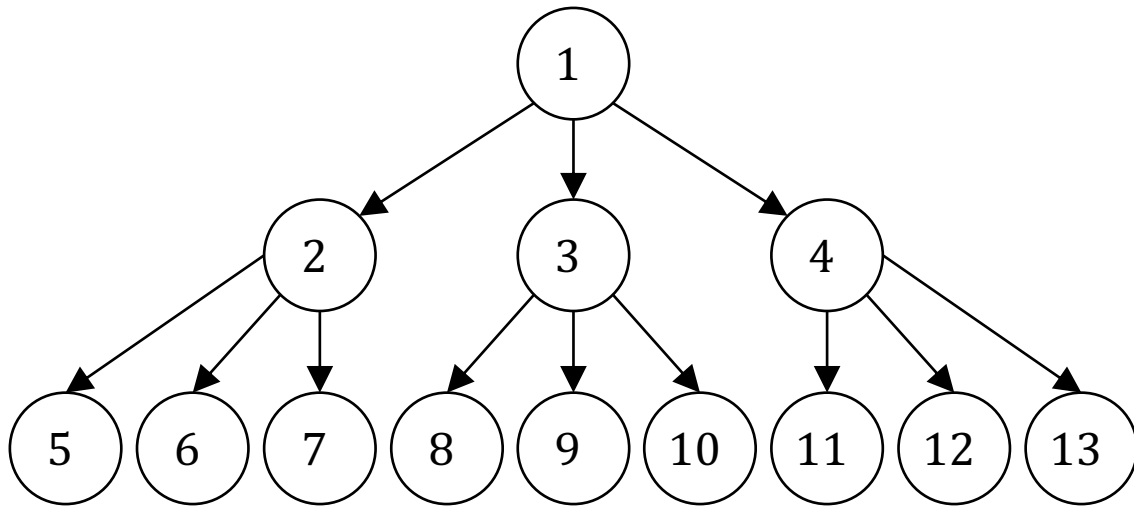
Verifying intervention sets

- Given a DAG G , we call a set \mathcal{J} of interventions a **verifying intervention set (VIS)** if the \mathcal{J} -essential graph of G has no undirected edges.

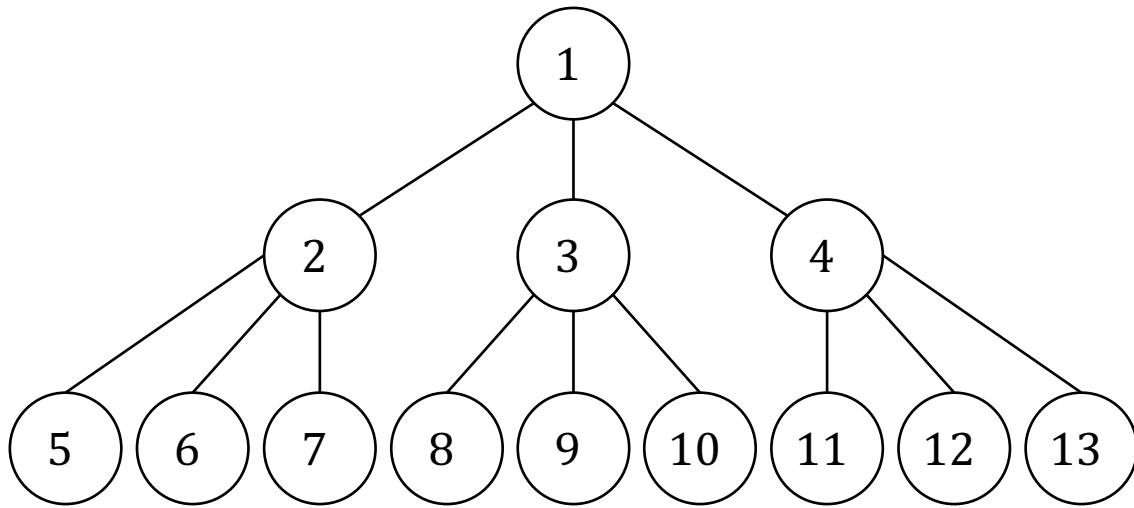
Verifying intervention sets

- Given a DAG G , we call a set \mathcal{J} of interventions a **verifying intervention set (VIS)** if the \mathcal{J} -essential graph of G has no undirected edges.
- We call a set \mathcal{J} of interventions a **minimal VIS (MVIS)** if it has the minimum cardinality of all VISes.

Special case: trees

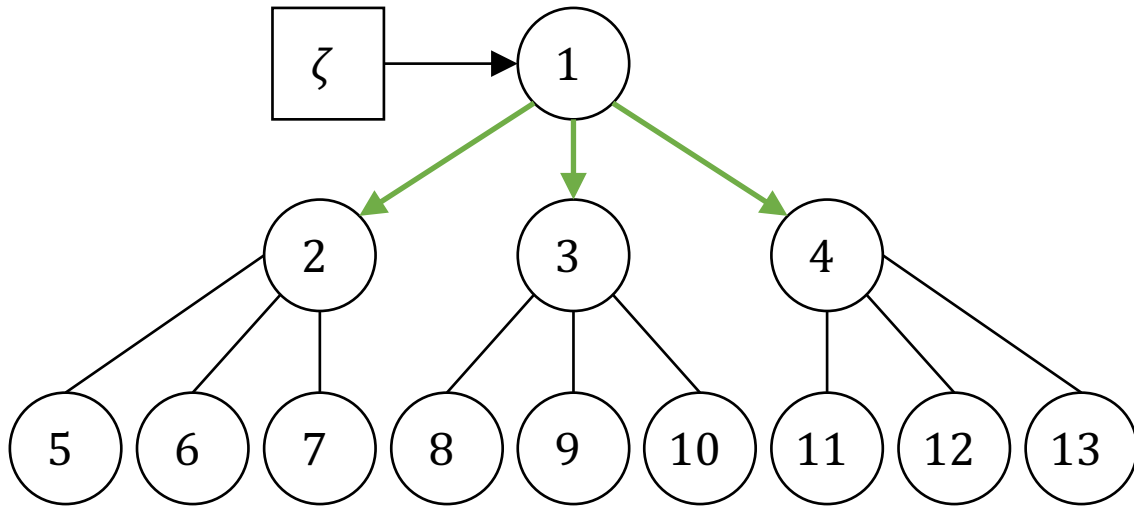


Special case: trees

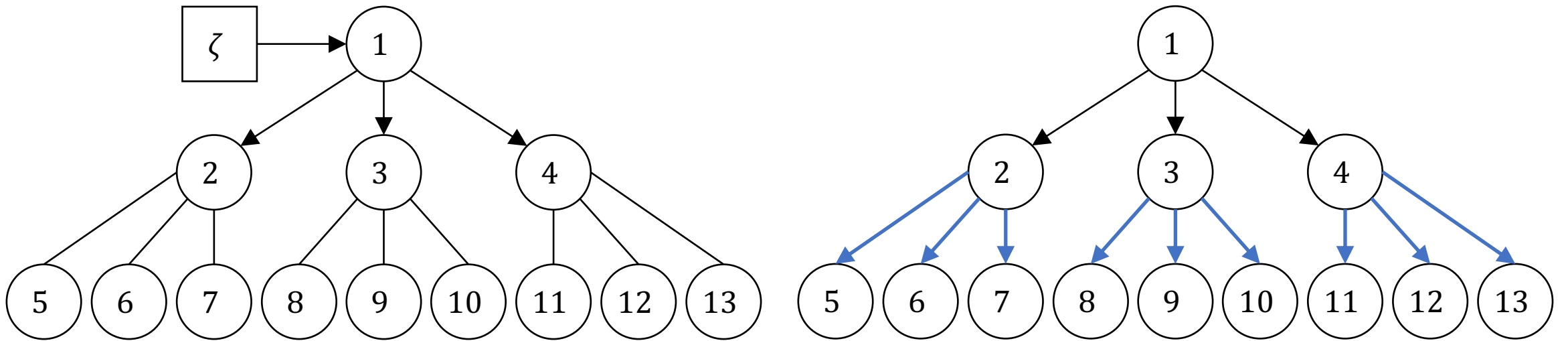


The essential graph

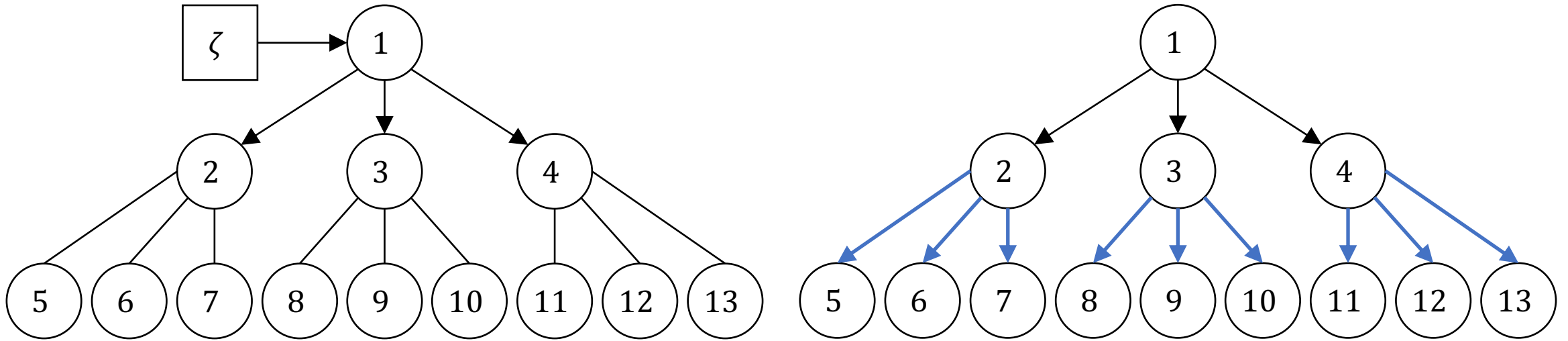
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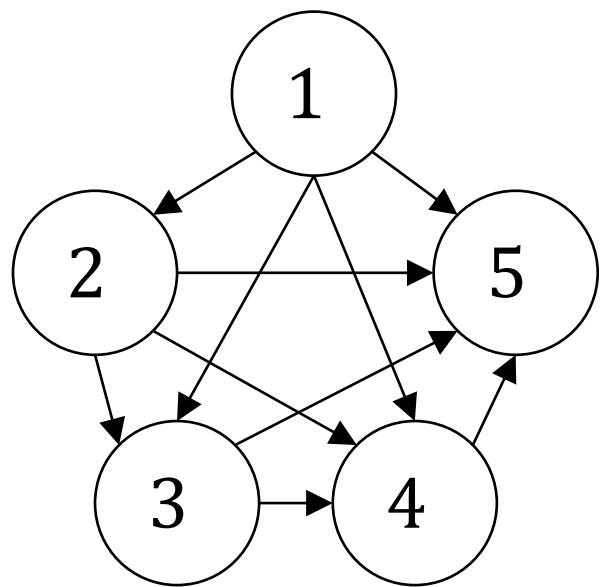


Special case: trees

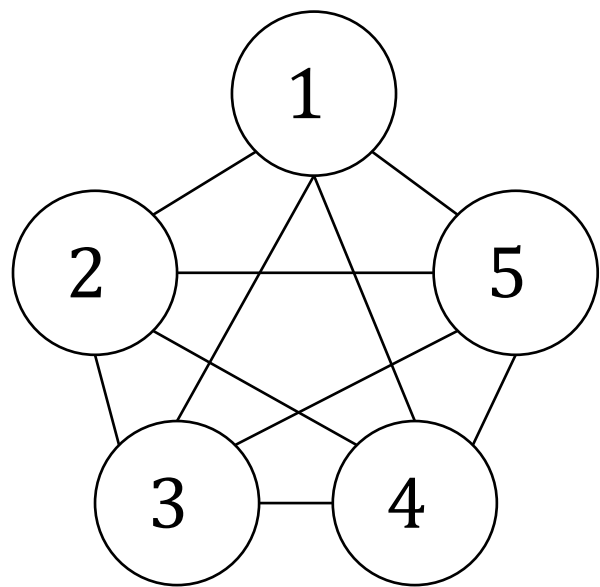


Only **one** intervention
is required

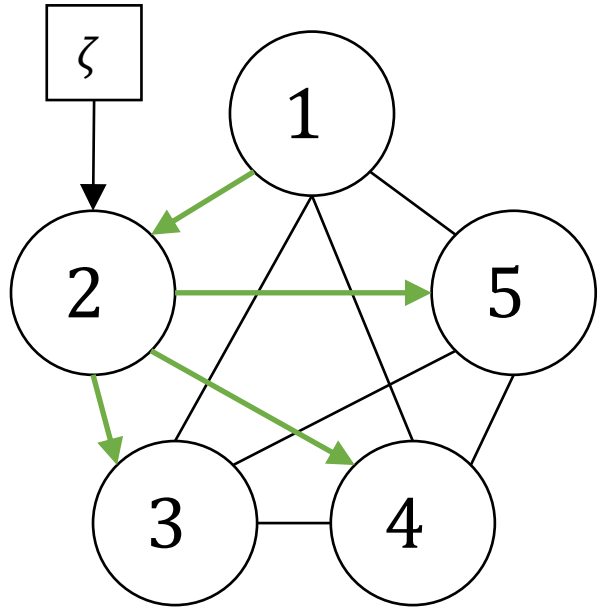
Special case: cliques



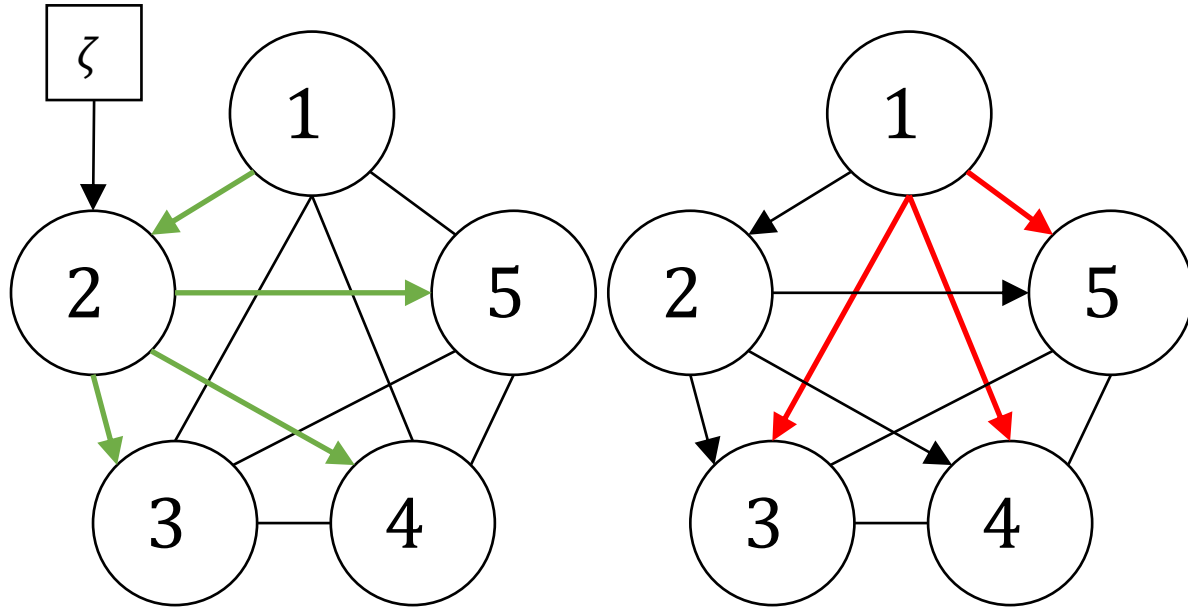
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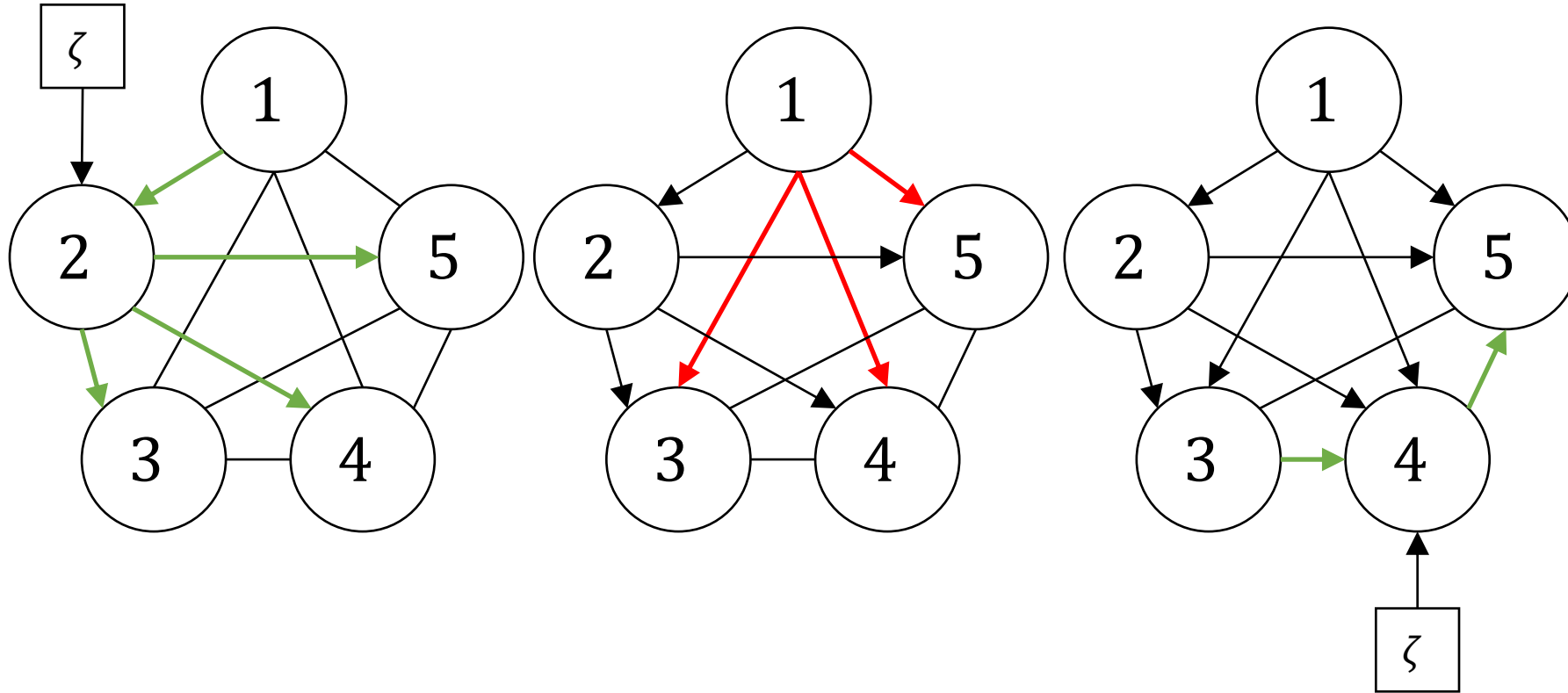
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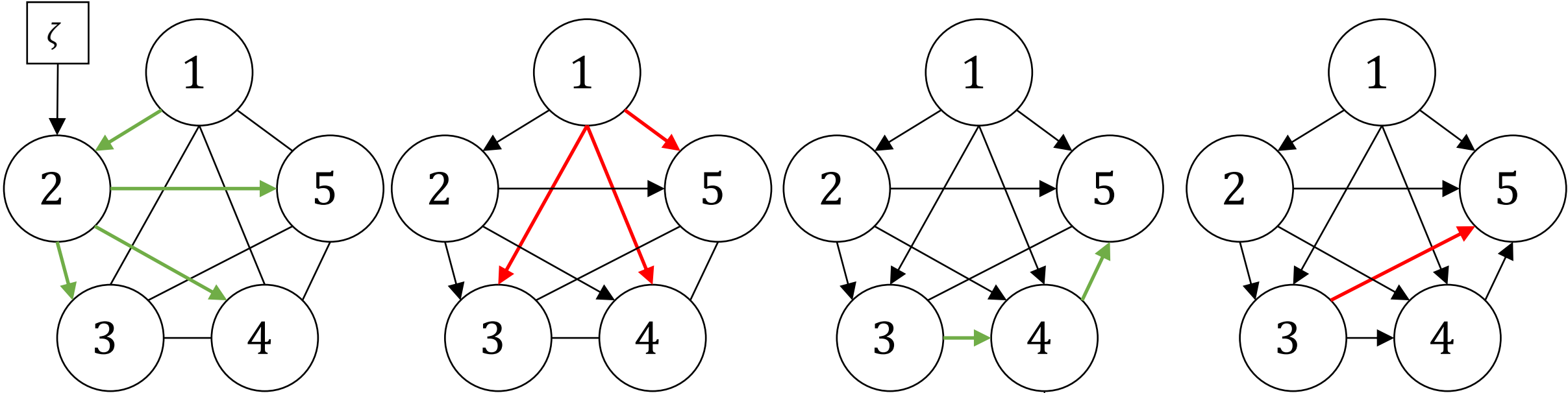
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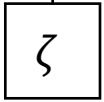
Special case: cliques



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Requires $\lfloor \frac{p}{2} \rfloor$
interventions for a
clique on p nodes

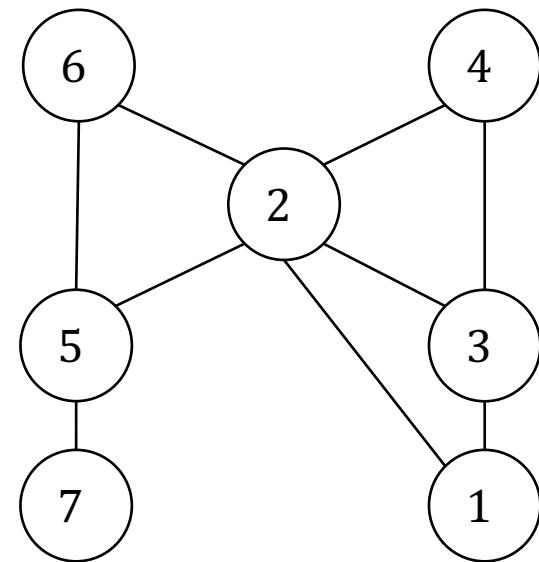


A general solution

Papers

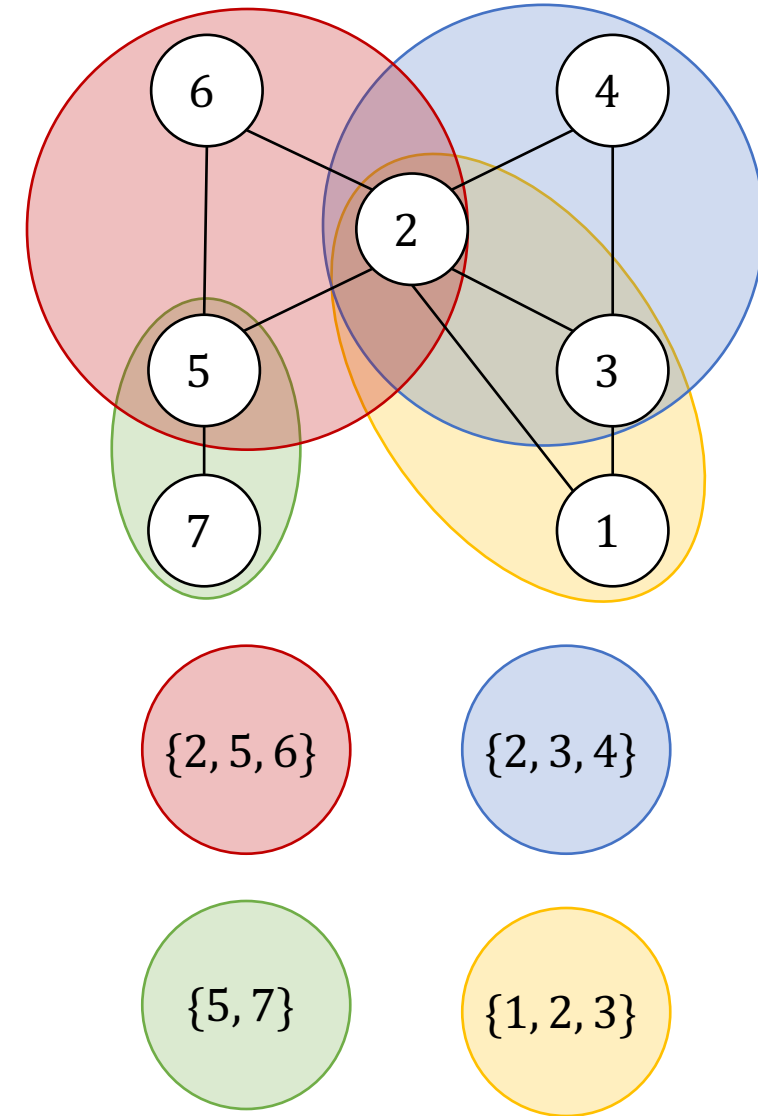
- **[SMG+]** *Active Structure Learning of Causal DAGs via Directed Clique Trees*, **Squires, C.**, Magliacane, S., Greenewald, K., Katz, D., Kocaoglu, M. and Shanmugam, K., 2020.

Clique trees of chordal graphs



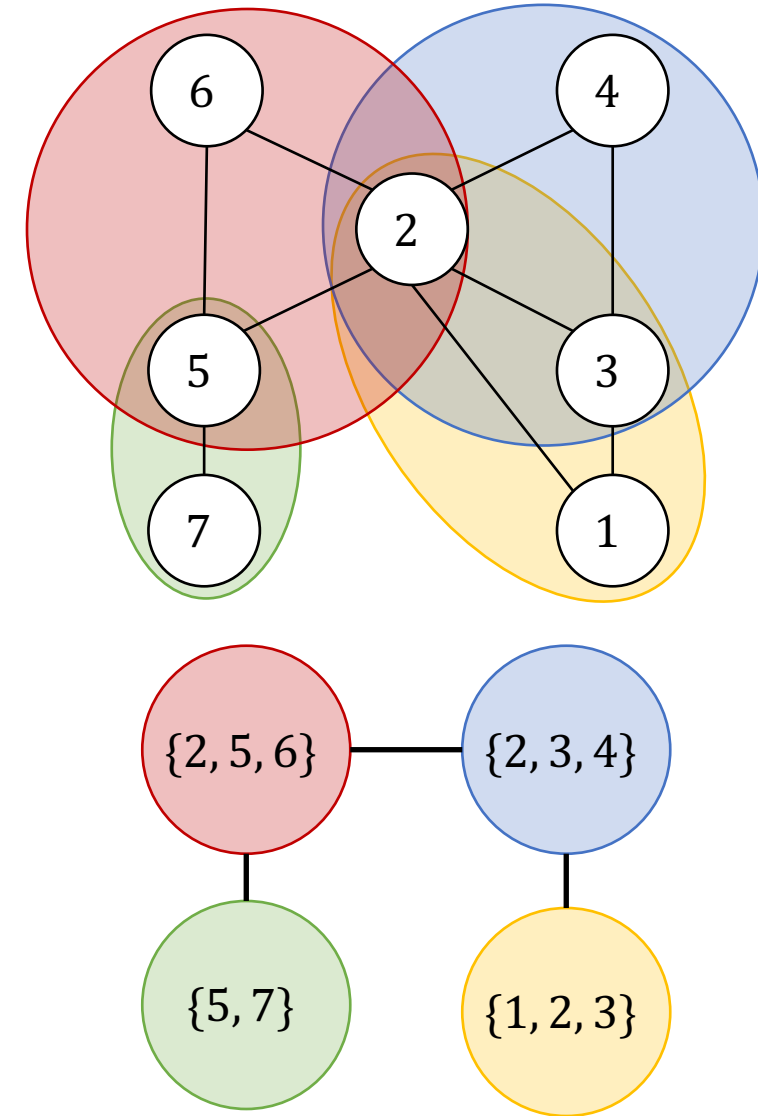
Clique trees of chordal graphs

- A new graph with:
 - Nodes being the **maximal cliques** of the original graph.



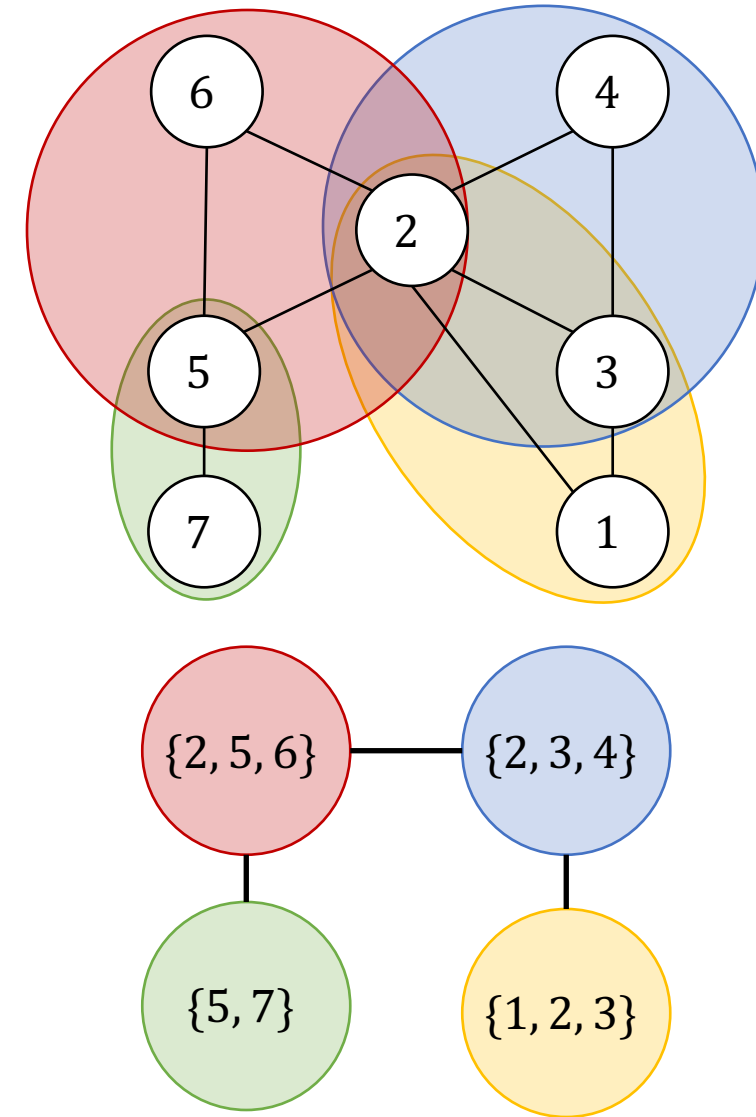
Clique trees of chordal graphs

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 - Edges forming a tree such that the **running intersection property** holds.

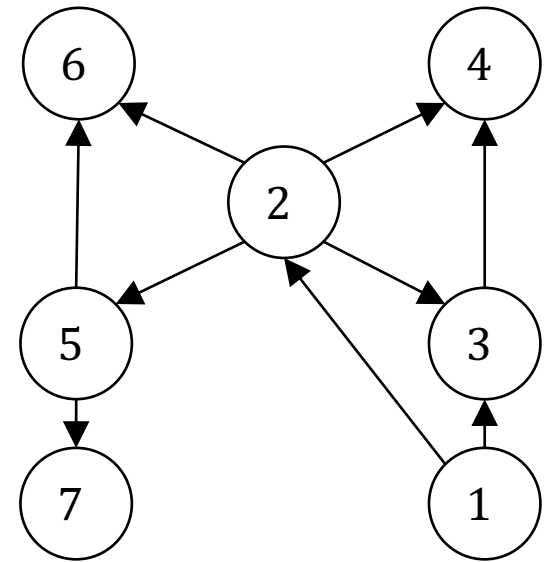


Clique trees of chordal graphs

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 - Edges forming a tree such that the **running intersection property** holds.
- The running intersection property: given two cliques C_1 and C_2 , their intersection $C_1 \cap C_2$ is contained in all cliques along the path from C_1 to C_2 .

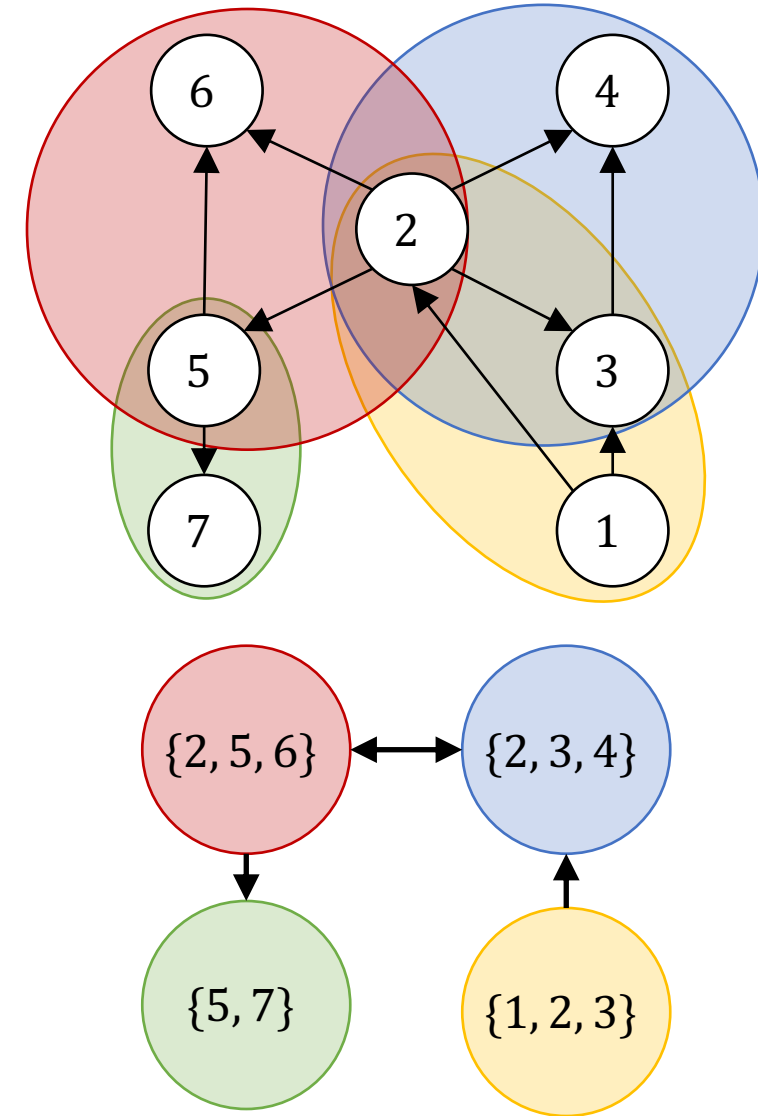


Directed Clique Trees (DCTs)



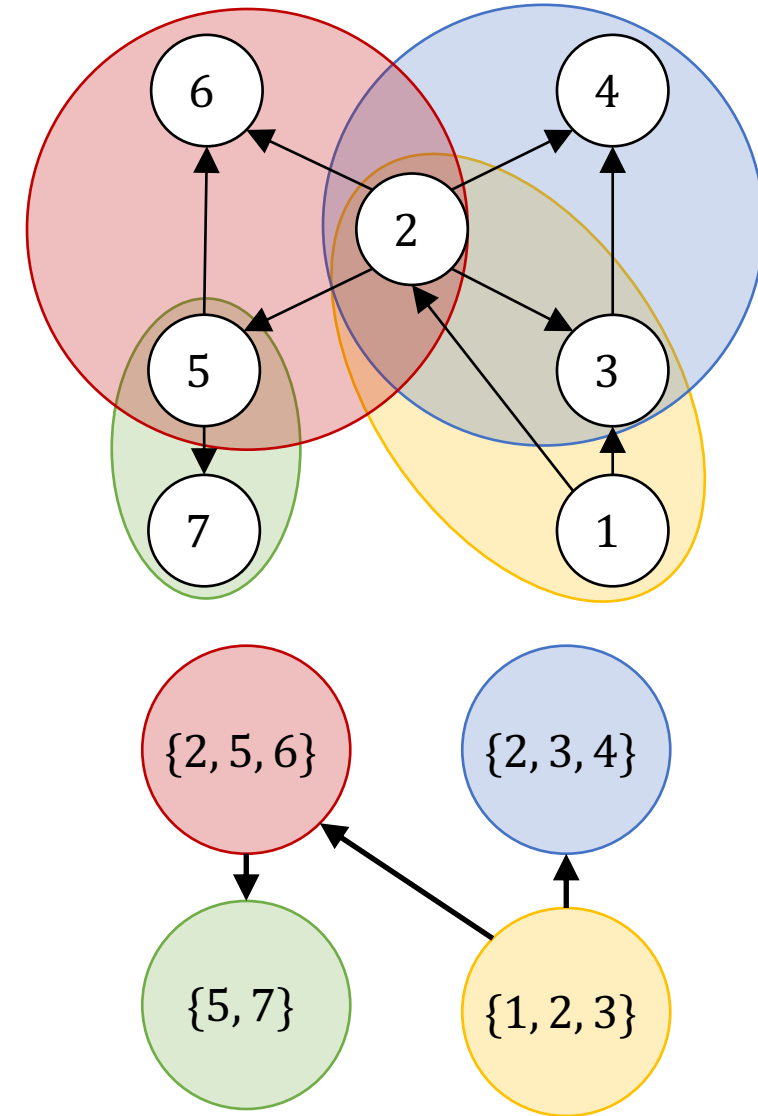
Directed Clique Trees (DCTs)

- Same vertices and adjacencies as a clique tree.
- Given adjacent cliques C_1 and C_2 , add an arrowhead at C_1 if:
 - for all $v_{12} \in C_1 \cap C_2$ and $v_2 \in C_2 \setminus C_1$, we have $v_{12} \rightarrow v_2$



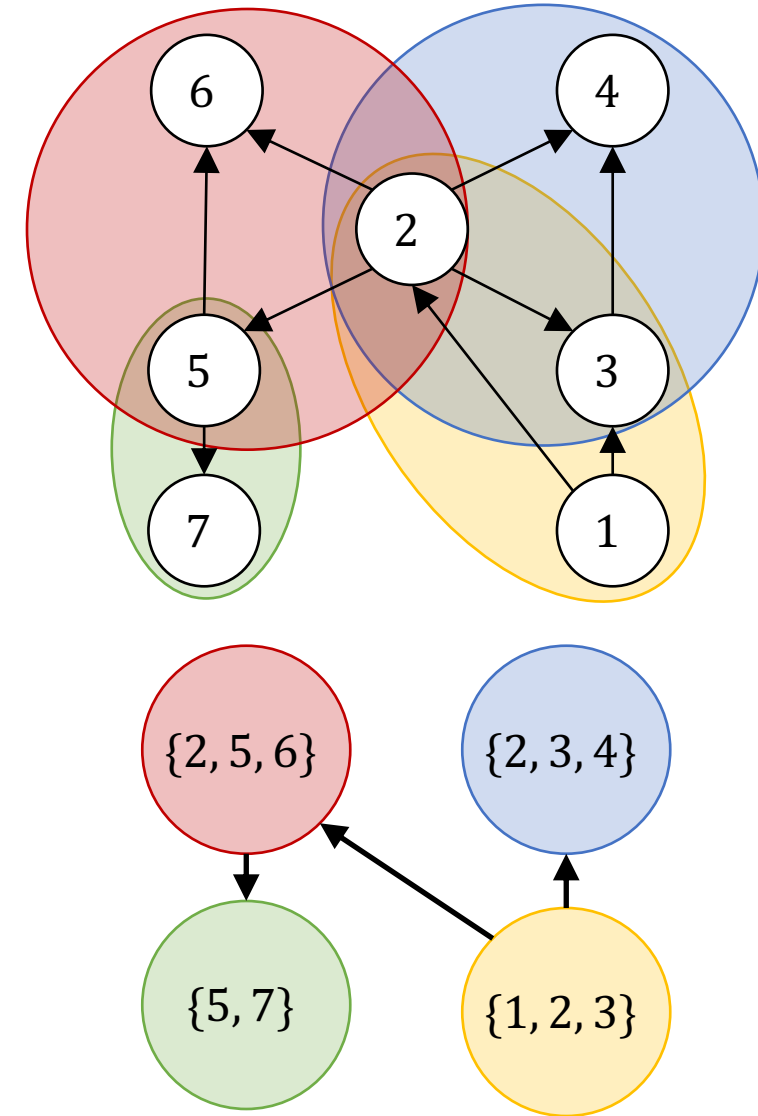
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- There can be multiple DCTs for a single graph.



Directed Clique Trees (DCTs)

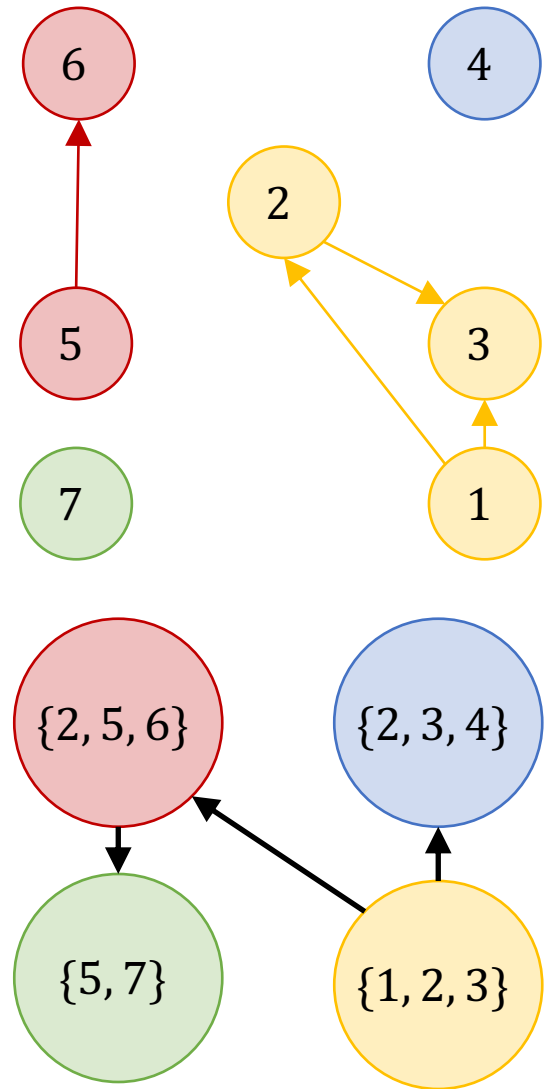
- There can be multiple DCTs for a single graph.
- We can always¹ find a DCT such that each clique has at most one parent (i.e., no colliders).



¹Requires a **contraction operation** that joins cliques connected by bidirected edges.

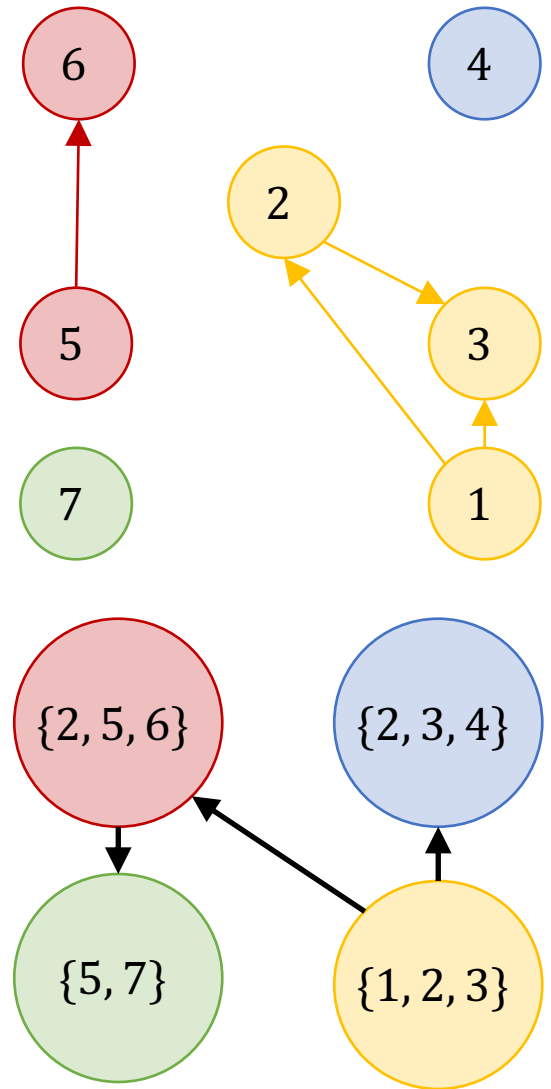
Residuals in DCTs

- The **residual** of a clique is the clique minus its parents.



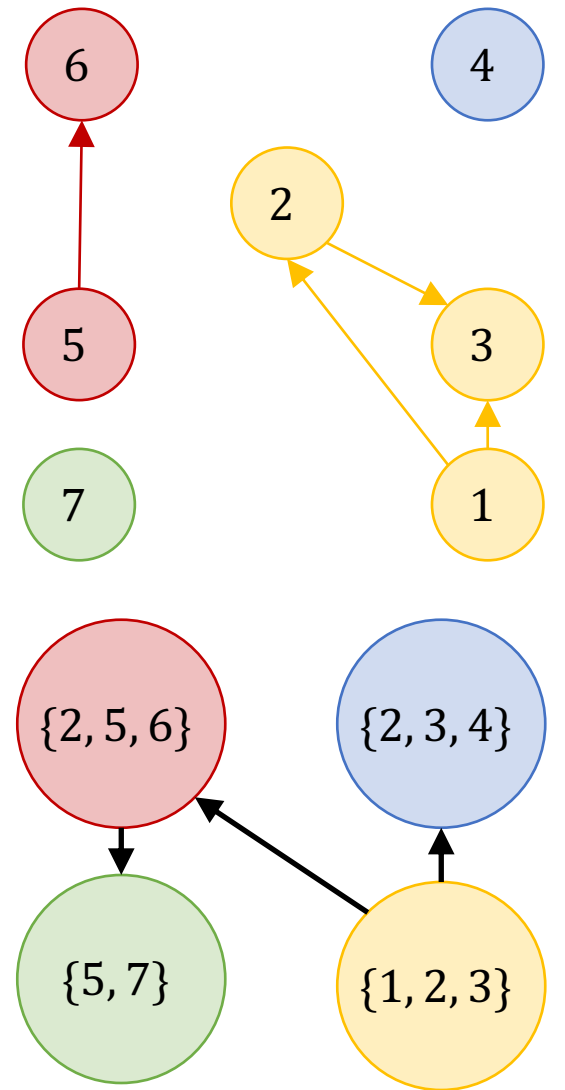
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Residuals in DCTs

- The **residual** of a clique is the clique minus its parents.
- Orientations in one residual are independent of the orientations in other residuals.
- Since the residuals are cliques (or other simple structures when we have to contract bidirected edges), we can efficiently compute the MVIS.



What else?

Instance-wise competitive bounds

- The **instance-wise competitive ratio (ic-ratio)** of an intervention policy on a DAG G is the (expected) number of interventions used by the policy, divided by the size of the MVIS.

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- We develop and analyze the **DCT-policy**, showing that the ic-ratio is logarithmic in the number of cliques for certain types of graphs.

Thanks!