# Introduction to Experimental Design in Causality

**Chandler Squires** 

# Agenda

Settings for experimental design in causal structure learning and preliminaries

9:30-10:30

Non-adaptive experimental design strategies

11-12

Adaptive experimental design strategies

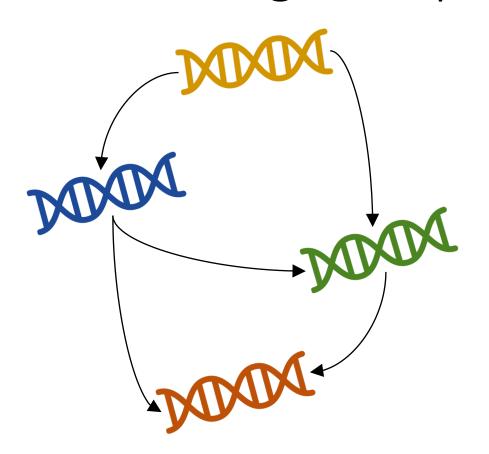
2-3

Targeted experimental design and other open challenges

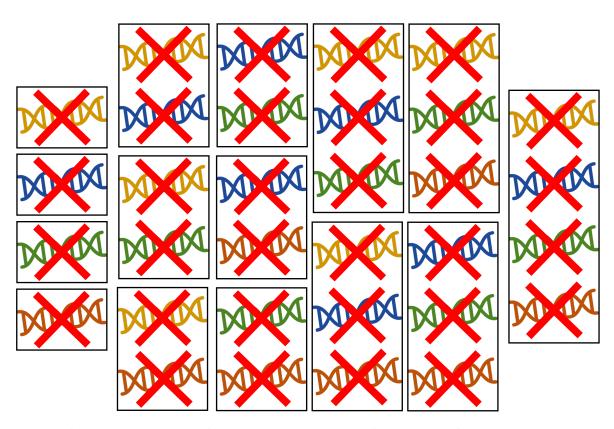
3:30-4:30

# Part I: Settings for experimental design in causal discovery and preliminaries

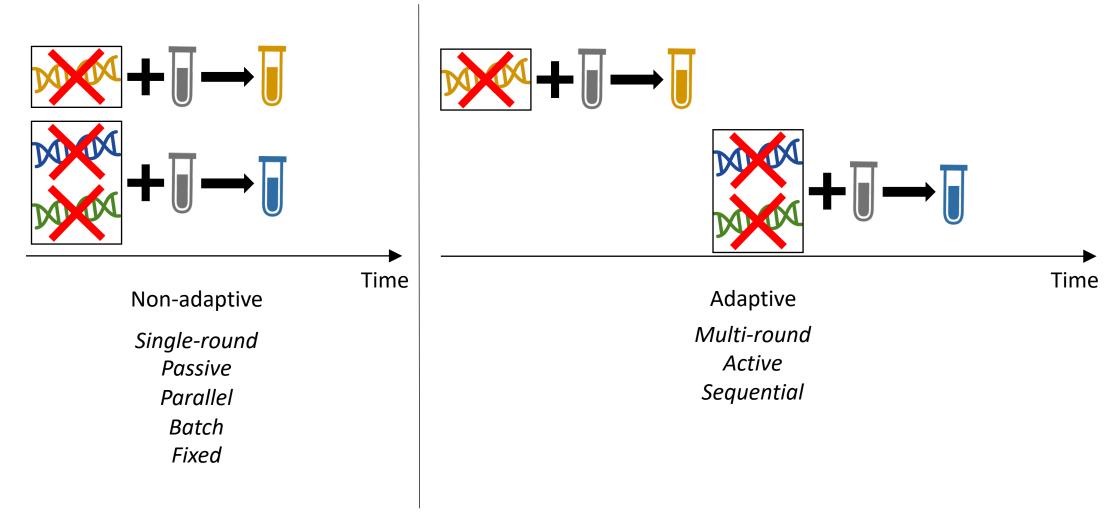
### Motivating Example: Genomics

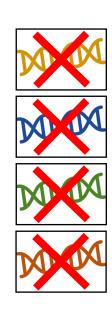


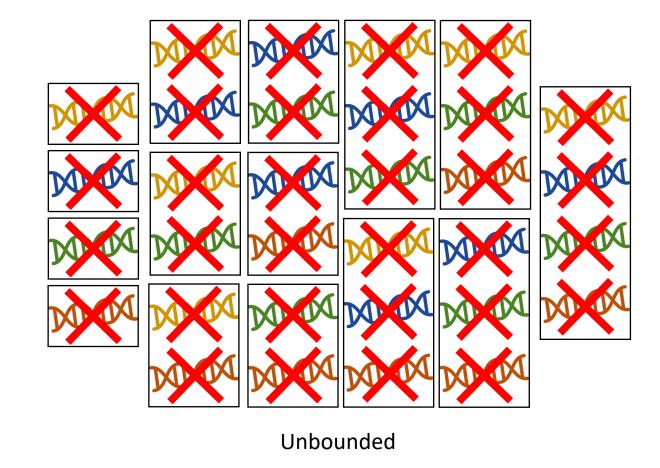
#### Knockout experiments:



How do we pick experiments to learn the underlying causal graph?







Bounded



max s.t. info(interventions) $cost(interventions) \le budget$ 

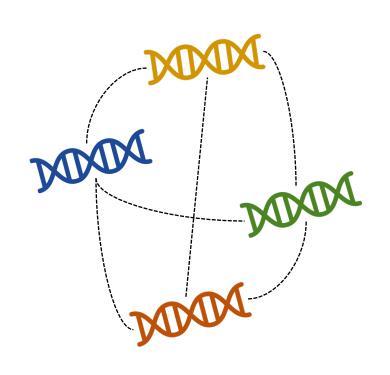
Fixed budget



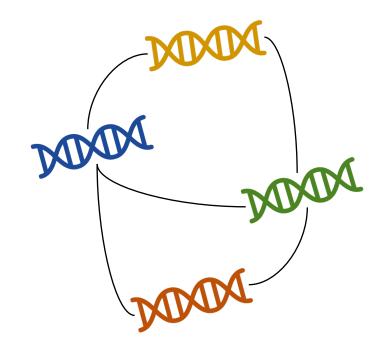
min s.t.

cost(interventions)
interventions identify G\*

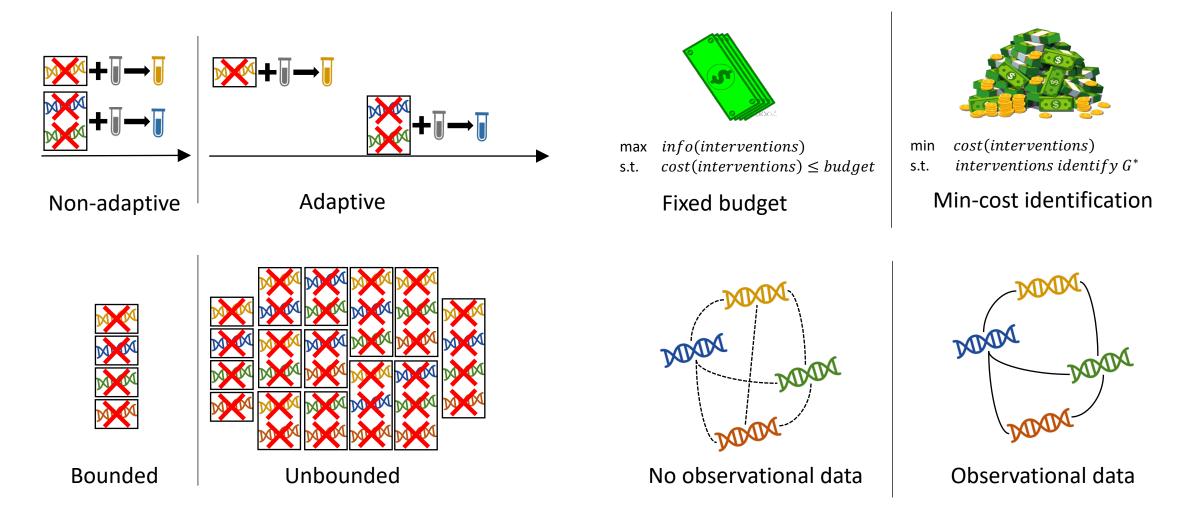
Minimum cost identification



No observational data



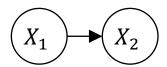
Observational data



# Additional assumptions for parts I, II, and III

- Noiseless setting: we obtain an infinite amount of data from each intervention, so that the only uncertainty is due to unidentifiability and not statistical noise.
  - i.e., if the distribution associated with an intervention  $I_k$  is  $f^k$ , then we assume access to *conditional invariance* oracle that tells us whether  $f^k(X_i \mid X_C) = f^{k'}(X_i \mid X_C)$  for any node i and set C
  - In practice, we can imagine setting a threshold  $\epsilon$  and obtaining enough interventional samples to distinguish any two distributions that are further than  $\epsilon$  under some measure (e.g., total variation distance)
- Causally sufficient setting: there are no unobserved confounders.

# Types of Interventions

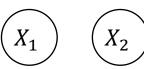


$$X_1 \leftarrow f_1(\varepsilon_1)$$
  
$$X_2 \leftarrow f_2(X_1, \varepsilon_2)$$





$$X_1 \leftarrow f_1(\varepsilon_1)$$
$$X_2 \leftarrow x_2$$



$$X_1 \leftarrow f_1(\varepsilon_1)$$
  
$$X_2 \leftarrow f_2'(\varepsilon_2')$$

$$X_1$$
  $X_2$ 

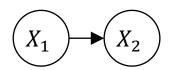
$$X_1 \leftarrow f_1(\varepsilon_1)$$
  
$$X_2 \leftarrow f_2'(X_1, \varepsilon_2')$$

Do-interventions 

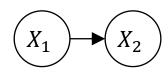
☐ Perfect interventions ☐ Soft interventions

aka: mechanism changes

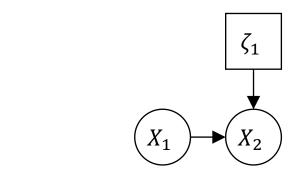
## Interventional Augmented Graphs



$$X_1 \leftarrow f_1(\varepsilon_1) X_2 \leftarrow f_2(X_1, \varepsilon_2)$$

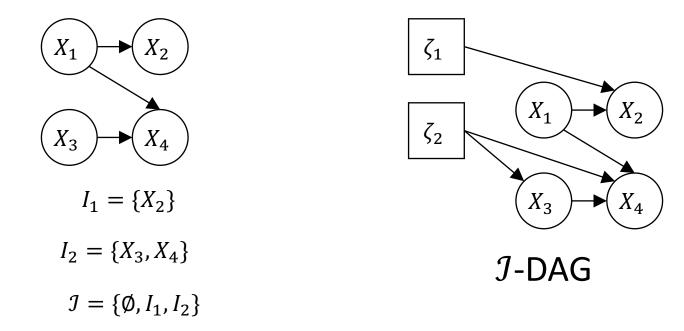


$$X_1 \leftarrow f_1(\varepsilon_1)$$
  
$$X_2 \leftarrow f_2'(X_1, \varepsilon_2')$$



$$X_1 \leftarrow f_1(\varepsilon_1)$$
  
$$X_2 \leftarrow (1 - \zeta_1) \cdot f_2(X_2, \varepsilon_2) + \zeta_1 \cdot f_2'(X_1, \varepsilon_2')$$

## Interventional Augmented Graphs



Given a DAG G and a set of interventions  $\mathcal{I} = \{I_1, \dots, I_K\}$ , the **interventional DAG** ( $\mathcal{I}$ -DAG), denoted  $G^{\mathcal{I}}$ , is obtained by introducing **intervention variables**  $\zeta_1, \dots, \zeta_K$  and edges  $\zeta_k \to X_i$  for  $X_i \in I_k$ 

Joint Causal Inference from Multiple Contexts (Mooij et al., 2020)

Characterizing and Learning Equivalence Classes of Causal DAGs under Interventions (Yang et al., 2018)

# Interventional Markov Equivalence

An indexed set of distribution  $(f^k)_{k=1}^K$  is **J-Markov** w.r.t. G if:

- each  $f^k$  is Markov w.r.t. G
- $f^k(X_i \mid X_{pa(i)}) = f^{k'}(X_i \mid X_{pa(i)})$  for all k, k' such that  $X_i \notin I_k \cup I_{k'}$ .

We call two DAGs  $G_1$  and  $G_2$   $\mathcal{J}$ -Markov equivalent if every  $(f_k)_{k=1}^K$  which is  $\mathcal{J}$ -Markov w.r.t.  $G_1$  is  $\mathcal{J}$ -Markov w.r.t.  $G_2$  and vice versa.

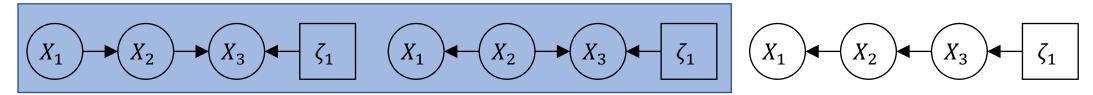
The  $\mathcal{I}$ -Markov equivalence class of G is denoted  $[G]_{\mathcal{I}}$ . The observational Markov equivalence class (i.e.,  $\mathcal{I} = \{\emptyset\}$ ) is denoted [G].

# Characterization of Interventional Markov Equivalence

Theorem (Hauser and Bühlmann, 2012, Yang et al., 2018)

Two DAGs  $G_1$  and  $G_2$  are  $\mathcal{I}$ -Markov equivalent iff. their  $\mathcal{I}$ -DAGs  $G_1^{\mathcal{I}}$  and  $G_2^{\mathcal{I}}$  have the same skeletons and v-structures.

Equivalently: Two DAGs are  $G_1$  and  $G_2$   $\mathcal{I}$ -Markov equivalent iff. they are Markov equivalent and when  $X_i \to X_j$  in  $G_1$  and  $|\{X_i, X_j\} \cap I_k| = 1$  for some  $I_k \in \mathcal{I}$ , then  $X_i \to X_j$  in  $G_2$ .

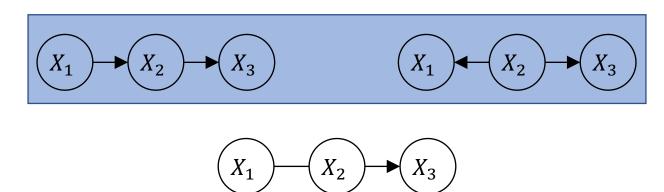


Characterization and Greedy Learning of Interventional Markov Equivalence Classes of Directed Acyclic Graphs (Hauser and Bühlmann, 2012)

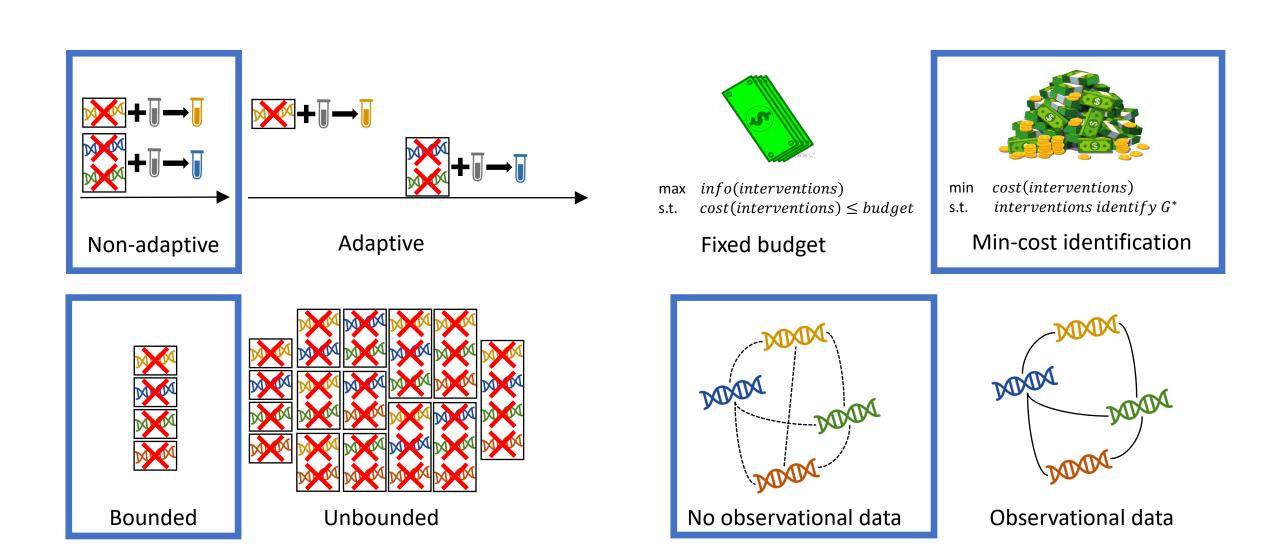
# $\mathcal{I}$ -essential graphs

Define the  $\mathcal{I}$ -essential graphs graph of  $G^*$ , denoted  $\mathcal{E}_{\mathcal{I}}(G^*)$ , as the mixed graph (directed and undirected edges) with:

- The same adjacencies as  $G^*$
- $X_i \to X_j$  in  $\mathcal{E}_{\mathcal{I}}(G^*)$  if  $X_i \to X_j$  for all  $G \in [G^*]_{\mathcal{I}}$
- $X_i X_j$  otherwise



# Part II: Non-adaptive experimental design strategies



#### Structural information from interventions

- If  $\{X_i, X_j\} \cap I_k = \{X_i\}$ , we call  $I_k$  an  $X_i$ -orientation test for  $(X_i, X_j)$
- If  $\{X_i, X_j\} \cap I_k = \emptyset$ , we call  $I_k$  an **adjacency test** for  $(X_i, X_j)$

#### Lemma (Eberhardt, 2008)

To determine the presence/absence and orientation of an edge between  $X_i$  and  $X_j$ , either of the following are sufficient and in the worst case necessary:

- 1. A structural orientation test and a structural adjacency test.
- 2. An  $X_i$ -orientation test and a  $X_i$ -orientation test.

# Single-node interventions

**Q:** How many single-node interventions are sufficient and worst-case necessary for finding any  $G^*$  on p nodes?

s.t.  $\forall G^*, \mathcal{I} identifies G^*$   $\forall I \in \mathcal{I}, |I| \leq 1$ 

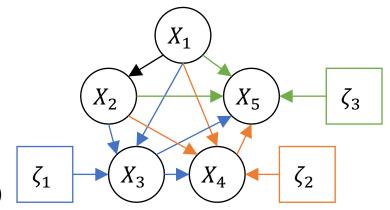
#### **Sufficiency**

All pairs of variables are subject to adjacency tests.

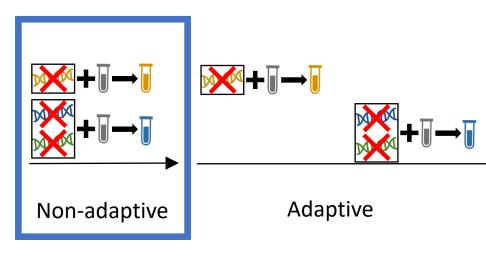
All pairs have at least one member subject to an orientation test.

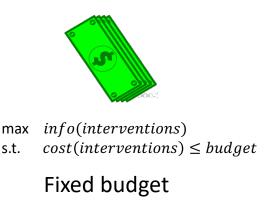
#### **Worst-case necessity**

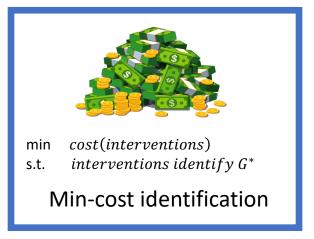
If we only intervene on p-2 variables, and the two we don't intervene on happen to be the most upstream.

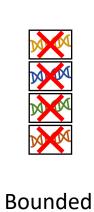


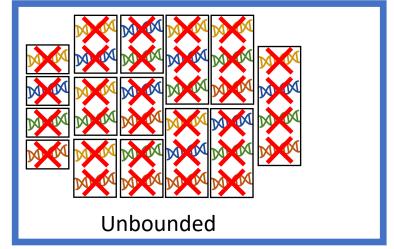
<sup>1</sup>Causation and Intervention (Eberhardt, 2008)

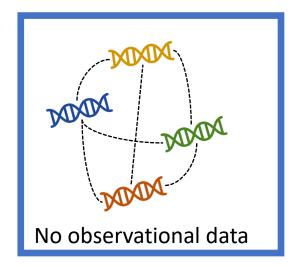


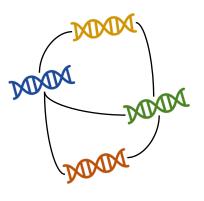












Observational data

#### Unbounded interventions

**Q:** How many (unrestricted) interventions are sufficient and worst-case necessary for finding any  $G^*$  on p nodes?

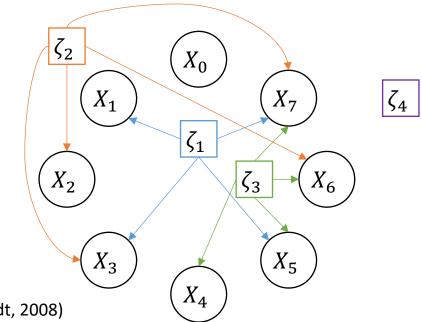
$$A^1: [\log_2 p] + 1$$

#### **Sufficiency**

Let b(k) denote the binary representation of an integer k, and  $b(k)_j$  the j-th rightmost digit in this representation.

$$b(6) = 110, b(6)_1 = 0, b(6)_2 = 1, \text{ etc.}$$

Label the vertices 0, ..., p-1 and let  $I_j = \{k \mid b(k)_j = 1\}$ , for  $j = 1, ..., \lfloor \log_2 p \rfloor + 1$ .



s.t.

 $\forall G^*, \mathcal{I} identifies G^*$ 

<sup>1</sup>Causation and Intervention (Eberhardt, 2008)

#### Unbounded interventions

**Q:** How many (unrestricted) interventions are sufficient and worst-case necessary for finding any  $G^*$  on p nodes?

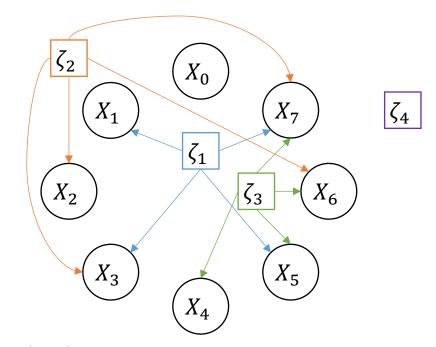
**A:**  $[\log_2 p] + 1$ 

#### **Sufficiency**

Each pair is subjected to an orientation test, since any two numbers differ in at least one position in their binary expansions.

If p is a power of 2, then there's an empty intervention which gives an adjacency test for all pairs.

Otherwise, we can use the fact that every binary expansion has at least one zero to show that every pair is subjected to either the opposing orientation test, or an adjacency test.



s.t.

 $\forall G^*, \mathcal{I} identifies G^*$ 

#### Unbounded interventions

**Q:** How many (unrestricted) interventions are sufficient and worst-case necessary for finding any  $G^*$  on p nodes?

St.  $\forall G^*$ ,  $\mathcal{I}$  identifies  $G^*$ 

**A:**  $[\log_2 p] + 1$ 

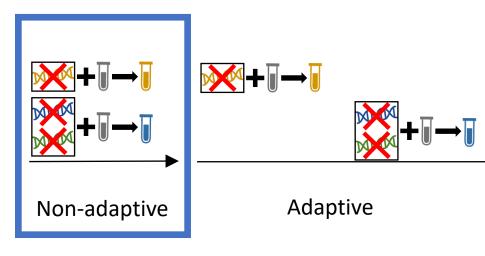
#### **Worst-case necessity**

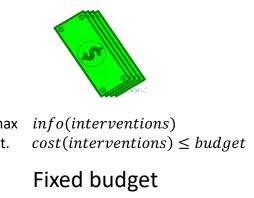
(induction) Assume the result holds for all q < p. Base cases on 2, 3, and 4 nodes can be done by hand.

Consider an intervention  $I_1$  on any number  $K \leq p$  nodes.

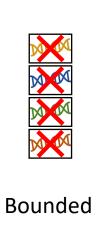
In the worst case, all p-K non-intervened nodes are upstream of these K nodes and form a complete graph, and there is a complete graph over the K intervened nodes as well.

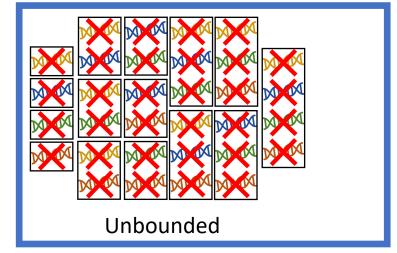
By the induction hypothesis, one of the remaining graphs requires at least  $\left[\log_2\frac{p}{2}\right]+1=\left[\log_2p\right]$  interventions.

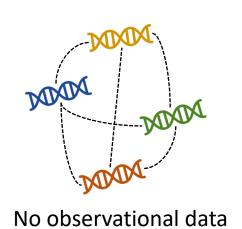


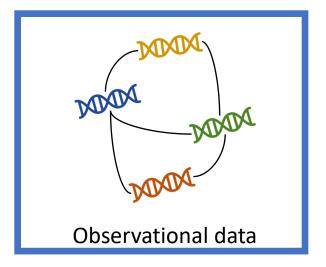












## Interventional Essential Graphs

The interventional essential graph is a *chain* graph with *chordal* chain components.

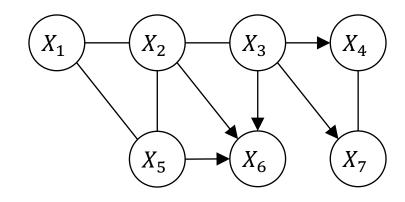
Furthermore, the orientations in each chain component are logically independent of one another.

# Chain graphs

A partially directed cycle in a mixed graph (i.e., a graph with both directed and undirected edges) is a sequence of distinct vertices  $v_0, v_1, ..., v_n$  such that:

- for each  $v_i$ , either  $v_{i-1}-v_i$  or  $v_{i-1} \rightarrow v_i$ , where  $v_{-1}=v_n$
- at least one edge in the path is directed

A **chain graph** is a mixed graph with no partially directed cycles.

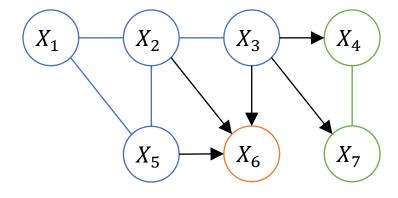


# Chain graphs

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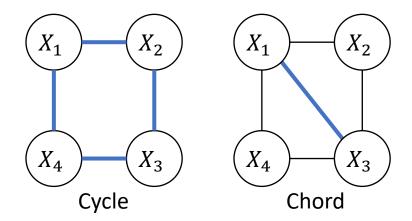
A **chain graph** is a mixed graph with no partially directed cycles.



The **chain components** of a chain graph are the connected components of the subgraph containing only the undirected edges.

# Chordal graphs

Given an undirected cycle, a **chord** is an edge that is not in the cycle but connects two nodes in the cycle.



A **chordal graph** is an undirected graph where all cycles of length  $\geq 4$  have a chord.

# The clique number of a graph

A **clique** in a graph is a set of nodes which are all adjacent.

The **clique number** of a graph G is the size of the largest clique, denoted  $\omega(G)$ .

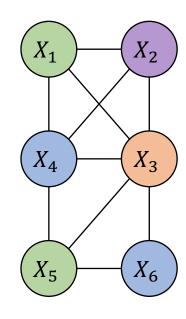
Note, this can be computed in linear time for a chordal graph G.

# Colorings of a graph

A **coloring** assigns a color to each node such that adjacent nodes do not have the same color.

The **chromatic number** of a graph is the smallest number of colors sufficient to form a coloring.

In a chordal graph G, the chromatic number is equal to the clique number  $\omega(G)$ , and an optimal coloring can be found in linear time.



#### Unbounded interventions for a fixed MEC

Q: How many (unrestricted) interventions are sufficient and worst-case

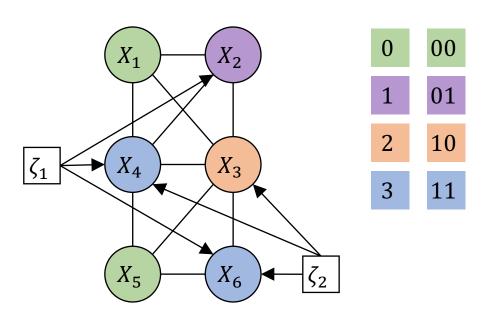
necessary for finding any  $G^* \in [G^*]$  on p nodes?  $\min_{\text{s.t.}} |\mathcal{I}|$   $\forall G^* \in [G^*]$ ,  $\mathcal{I}$  identifies  $G^*$ 

**A:**  $\lceil \log_2 \omega(\mathcal{E}(G^*)) \rceil$ 

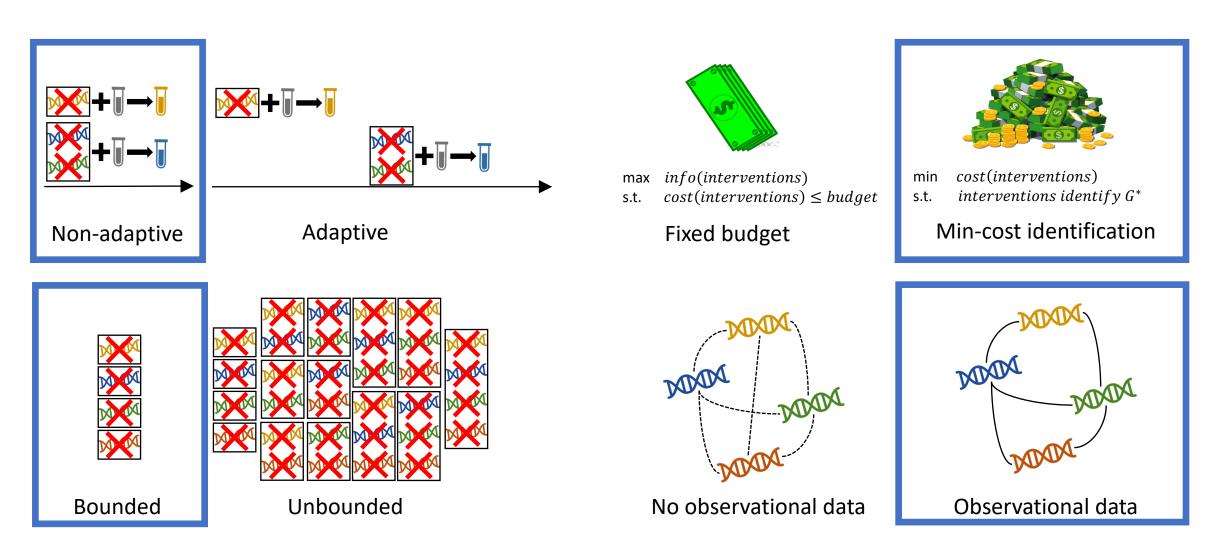
#### **Sufficiency**

Assign an optimal coloring to each chordal component, with each color corresponding to a number  $0, ..., \omega(\mathcal{E}(G^*))$ . Let c(j) denote the color (number) assigned to node j.

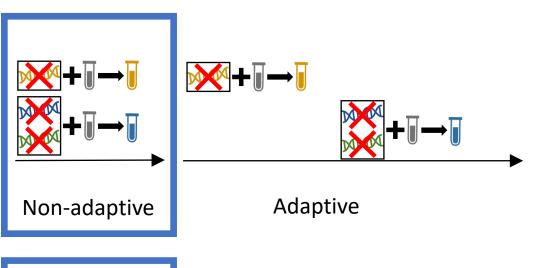
Then, let  $I_k = \{j \mid b(c(j))_k = 1\}$ . The argument follows the same as in the case with no observational data.

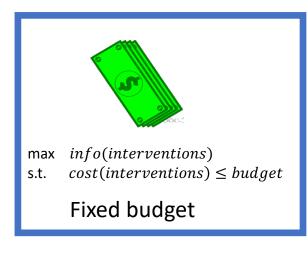


Two Optimal Strategies for Active Learning of Causal Models From Interventional Data (Hauser and Bühlmann, 2012)

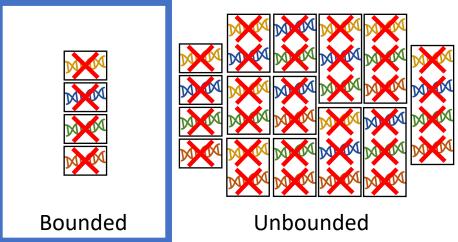


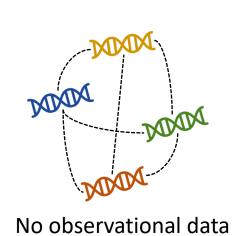
- Up to B nodes per intervention: <u>Learning Causal Graphs with Small Interventions</u> (Shanmugam et al., 2015)
- With different costs per intervention: Cost-Optimal Learning of Causal Graphs (Kocaoglu et al., 2017)

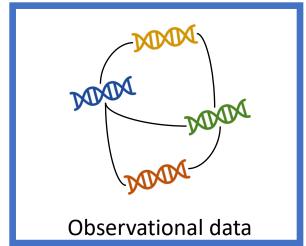












# Single-node interventions for a fixed MEC, with a fixed budget

**Task:** Pick K single-node interventions to maximize the expected number of oriented edges.

$$\max \quad \mathbb{E}_{G \sim Unif([G^*])}[dir(\mathcal{E}_{\mathcal{I}}(G))]$$
 s.t. 
$$|\mathcal{I}| \leq K$$
 
$$\forall I \in \mathcal{I}, |I| \leq 1$$

# Submodularity

- A set function  $f: 2^{\Omega} \to \mathbb{R}$  is **submodular** if for all  $A \subset B$  and  $b \in \Omega \setminus B$ , we have  $f(A \cup \{b\}) f(A) \ge f(B \cup \{b\}) f(B)$
- A set function is **monotonically increasing** if for all  $A \subset B$ , we have  $f(A) \leq f(B)$
- Greedy optimization (up to size *K*):
  - Start from  $A_0 = \emptyset$ , then repeat K times:
  - Let  $a_{i+1} = \operatorname{argmax}_{a \in \Omega \setminus A_i} f(A_i \cup \{a\})$ , and  $A_{i+1} = A_i \cup \{a_{i+1}\}$
- For a monotonically increasing submodular function, greedy optimization finds a set  $A_K$  such that  $f(A_K) \ge \left(1 \frac{1}{e}\right) \max_{|A| \le K} f(A)$

Theorem (Ghassami et al., 2018)

Let  $\Omega$  be the set of single-node interventions, let G be any DAG, and let  $f_G(\mathcal{I}) = dir(\mathcal{E}_{\mathcal{I}}(G))$ . Then  $f_G(\mathcal{I})$  is monotonically increasing and submodular.

Theorem (Ghassami et al., 2018)

Let  $E_{\mathcal{I}_1}$  be the set of edges oriented by the set  $\mathcal{I}_1$  of single-node interventions, and  $E_{\mathcal{I}_2}$  the set of edges oriented by the set  $\mathcal{I}_2$  of single-node interventions. Then  $E=E_{\mathcal{I}_1}\cup E_{\mathcal{I}_2}$  is complete under Meek's orientation rules.

Why does this imply submodularity?

$$f_D(\mathcal{I} \cup \{i\}) - f_D(\mathcal{I}) = |E_{\{i\}} \setminus E_{\mathcal{I}}|$$

So, checking submodularity reduces to checking that  $\mathcal{I} \subset \mathcal{I}' \Rightarrow \left| E_{\{i\}} \setminus E_{\mathcal{I}} \right| \geq \left| E_{\{i\}} \setminus E_{\mathcal{I}'} \right|$ 

Theorem (Ghassami et al., 2018)

Let  $E_{\mathcal{I}_1}$  be the set of edges oriented by the set  $\mathcal{I}_1$  of single-node interventions, and  $E_{\mathcal{I}_2}$  the set of edges oriented by the set  $\mathcal{I}_2$  of single-node interventions. Then  $E=E_{\mathcal{I}_1}\cup E_{\mathcal{I}_2}$  is complete under Meek's orientation rules.

#### Proof sketch:

- Check the pre-conditions for each Meek rule which involves 2 oriented edges  $X_1 o X_2$  and  $X_3 o X_4$
- If E is not complete, then  $X_1 \to X_2$  must be from  $E_{\mathcal{I}_1}$  and  $X_3 \to X_4$  must be from  $E_{\mathcal{I}_2}$
- Show that  $X_1 \to X_2 \in E_{\mathcal{I}_1}$  implies that  $X_3 \to X_4 \in E_{\mathcal{I}_2}$

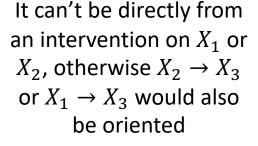
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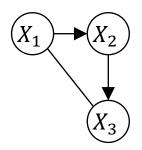
#### **Example:**

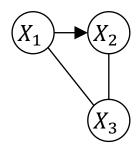
Say the "no cycles" rule is invoked

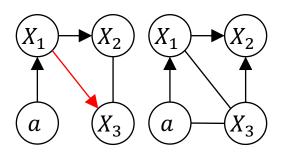
Can we have  $X_1 \rightarrow X_2 \in E_{\mathcal{I}_1}$  without any of the other edges?



It can't be from the "no new v-structures" rule, or else  $X_1 \rightarrow X_3$  or  $X_3 \rightarrow X_2$  would be oriented







# Counting and sampling from a Markov equivalence class

Theorem 4, Wienöbst et al., 2020

Let  $\mathcal{C}(G)$  denote the set of maximal cliques in a chordal graph G. Then, we can sample uniformly from [G] with p nodes and e edges in  $\mathcal{O}(p+e)$  with  $\mathcal{O}(|\mathcal{C}(G)|^2 \cdot p \cdot e)$  setup time.

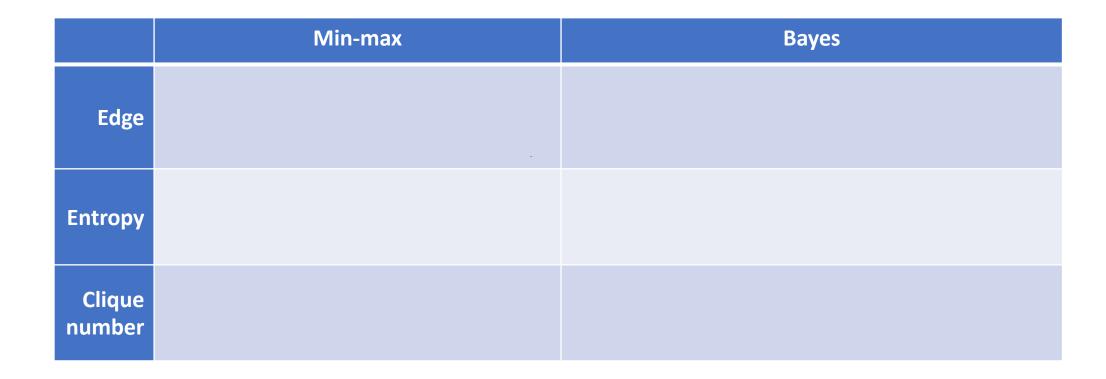
Thus, we may efficiently approximate  $\mathbb{E}_{G \sim Unif([G^*])}[dir(\mathcal{E}_{\mathcal{I}}(G))]$  for any  $\mathcal{I}$  by sampling and Monte-Carlo averaging.

Submodularity and monotonicity are preserved under positive linear combination, so the full problem is submodular.

Polynomial-Time Algorithms for Counting and Sampling Markov Equivalent DAGs (Wienöbst et al., 2020)

# Part III: Adaptive experimental design strategies

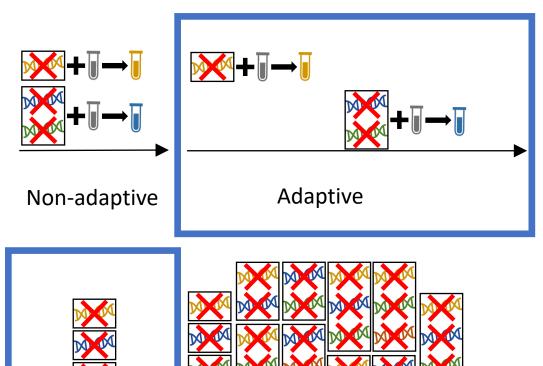
#### Greedy strategies

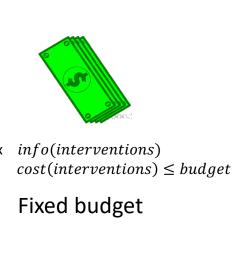


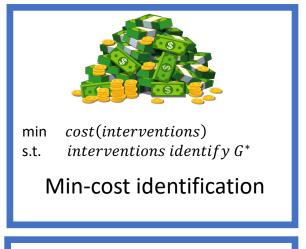
<sup>1</sup>Two Optimal Strategies for Active Learning of Causal Models From Interventional Data (Hauser and Bühlmann, 2012)

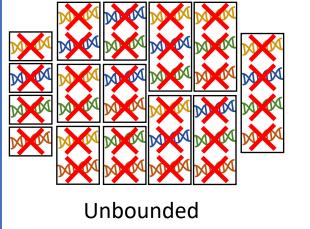
<sup>2</sup>Learning Causal Graphs with Small Interventions (Shanmugam et al., 2015)

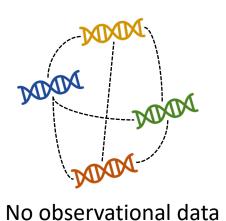
<sup>3</sup>Active Learning of Causal Networks with Intervention Experiments and Optimal Designs (He and Geng, 2008)

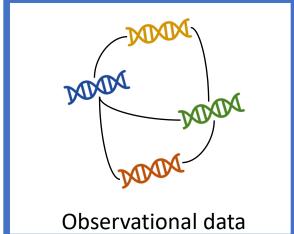




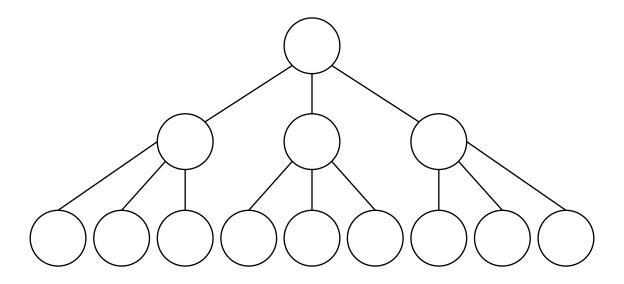




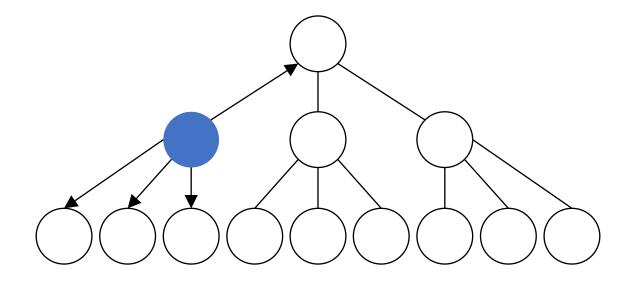


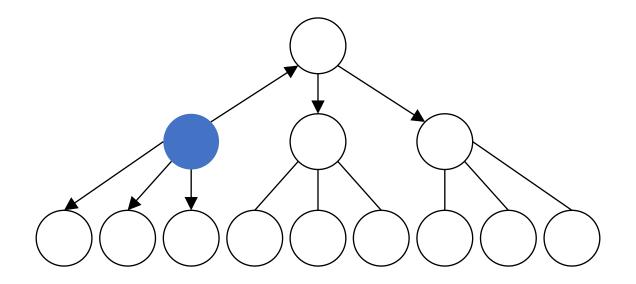


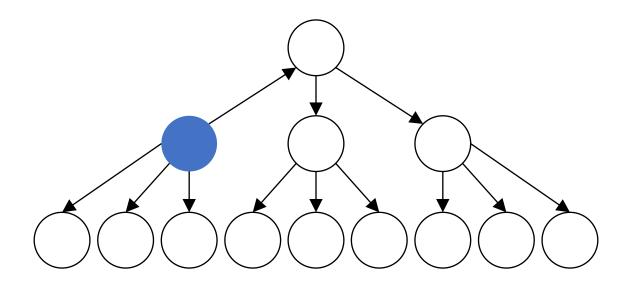
**Bounded** 



Essential graph

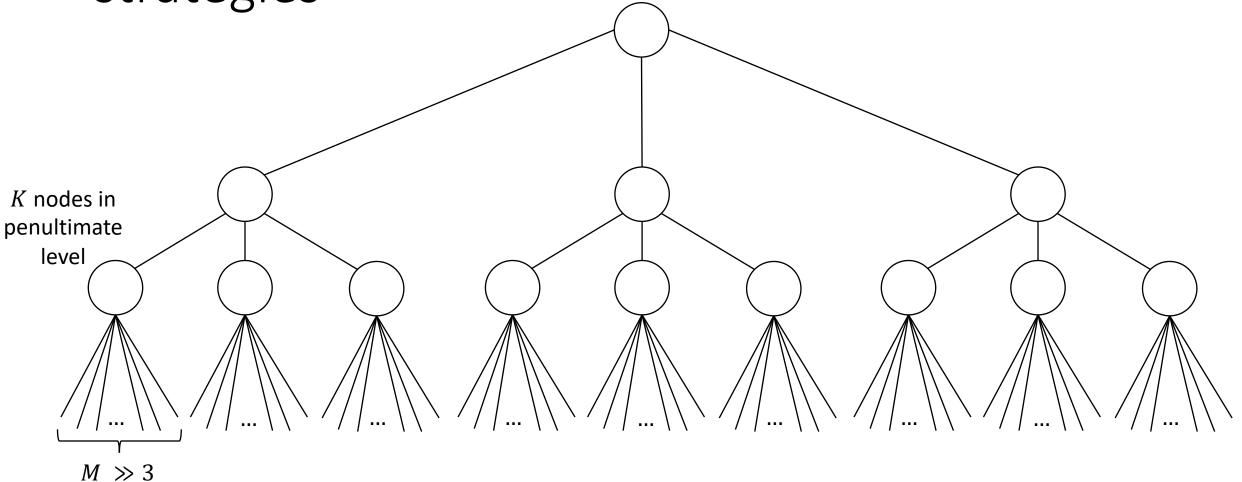






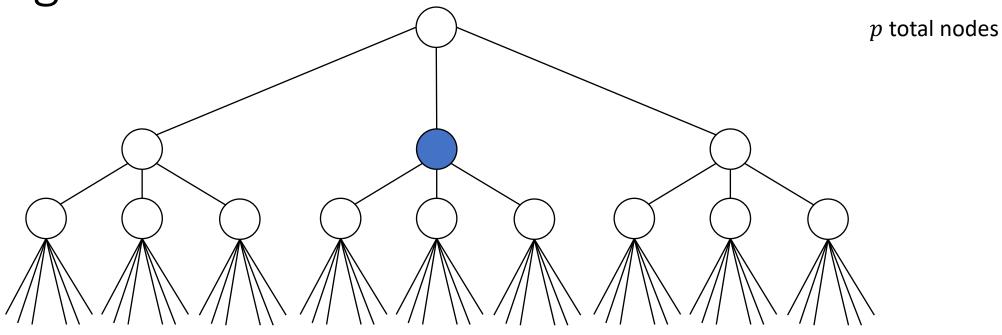
The root node uniquely determines the DAG

Suboptimality of information-greedy strategies



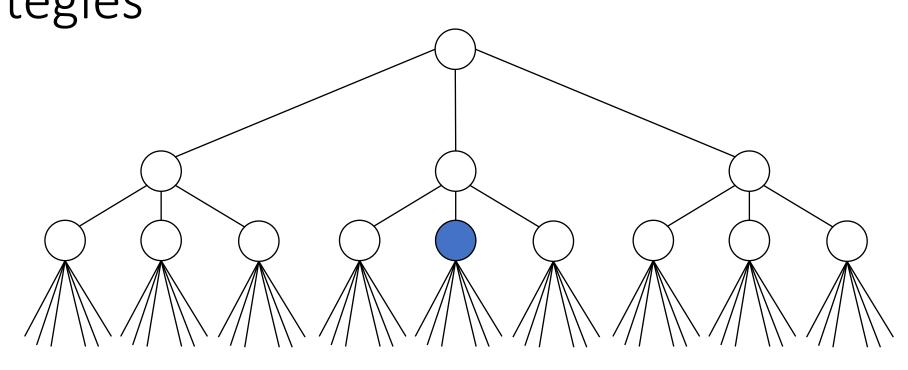
Suboptimality of information-greedy

strategies



Expected information gain 
$$\leq \frac{1}{p} \log_2 p + \frac{p-1}{p} \log_2 4 \approx 2$$

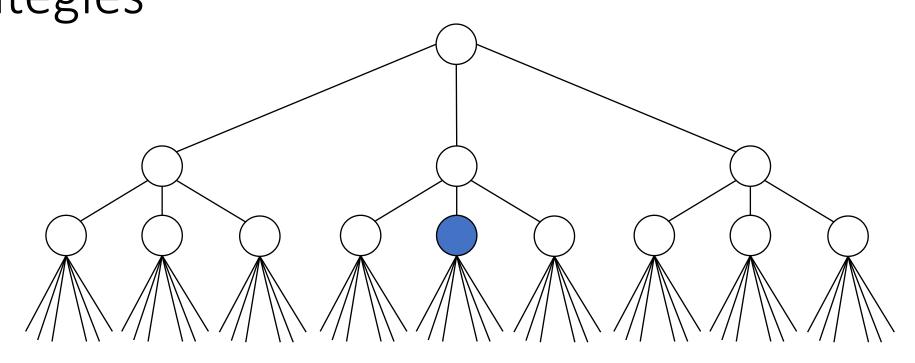
Suboptimality of information-greedy strategies



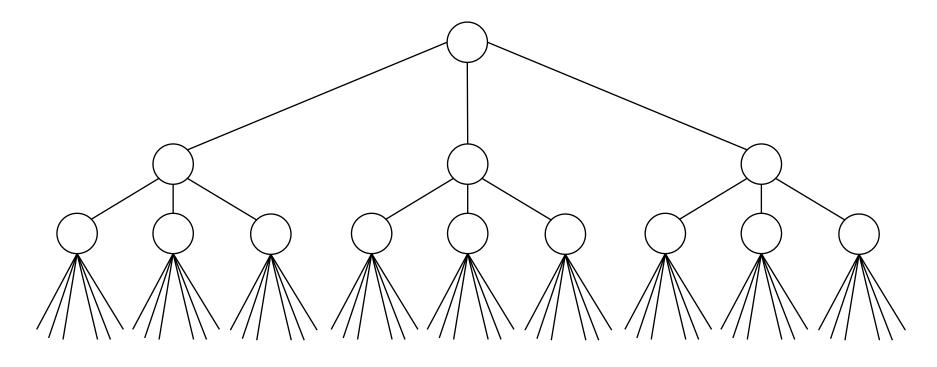
Expected information gain 
$$\geq \frac{1}{K} \log_2 p \geq \frac{1}{K} \log_2 M$$

M can be picked so that the nodes in the penultimate layer are always preferable (e.g.,  $M = O(2^{K^2})$ )

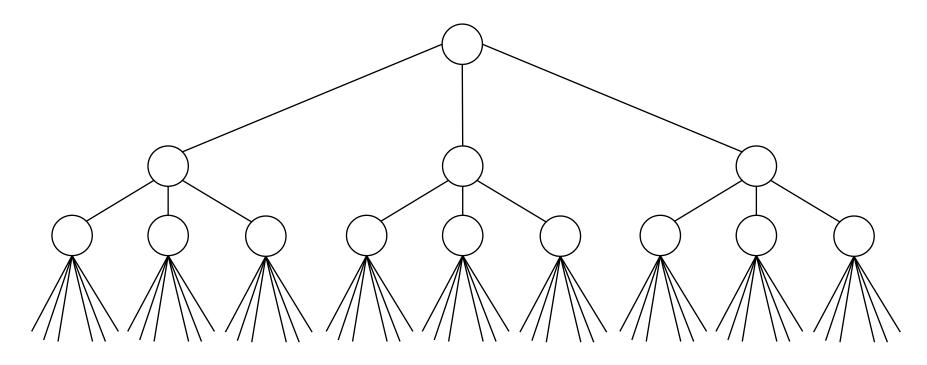
Suboptimality of information-greedy strategies



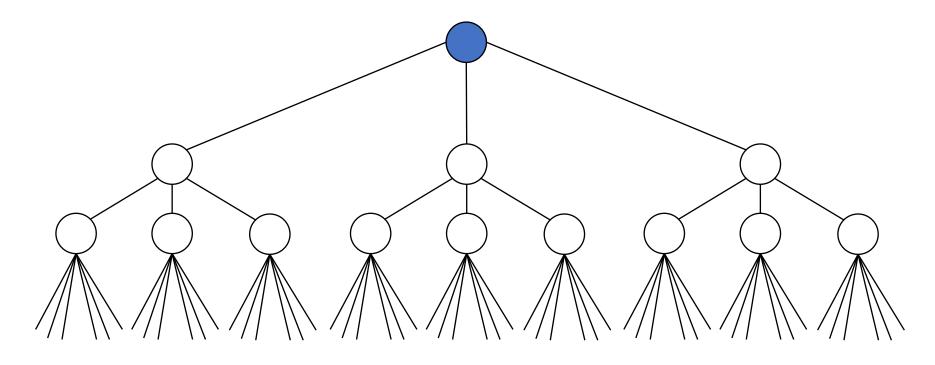
The expected number of interventions required by this strategy is at least K/2

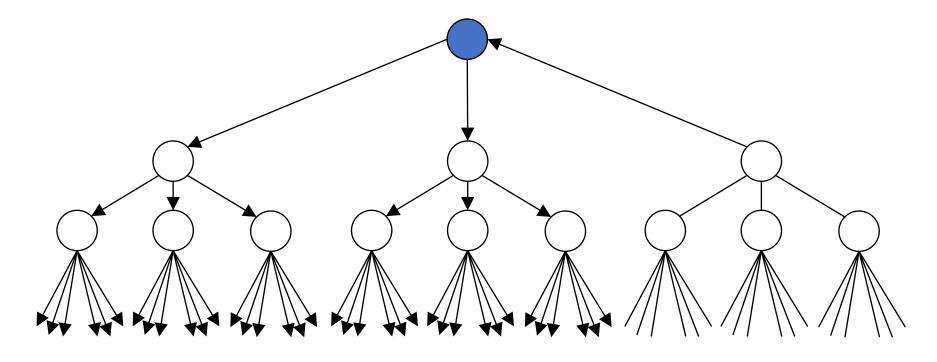


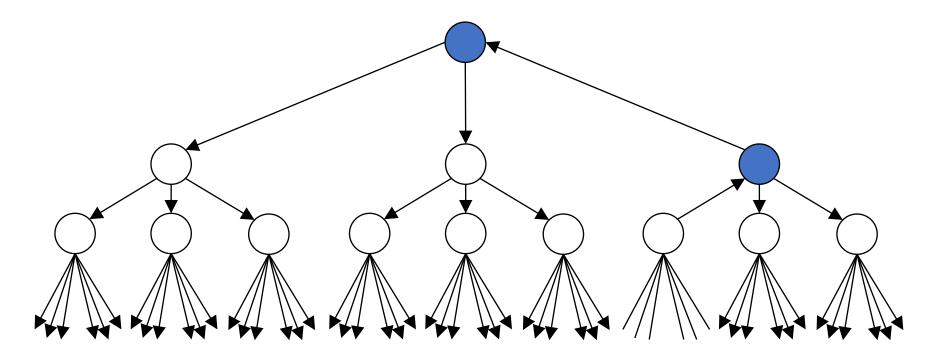
Central node algorithm<sup>1</sup> can find the root in at most  $(\log_3 K) + 1$  interventions



**Central node algorithm:** Intervene on a node such that, after removing it, the largest remaining connected component is as small as possible. Equivalent to min-max entropy choice.







- In a tree on p nodes, there is at least one node such that its removal splits the tree into components each having at most  $\frac{p}{2}$  nodes.<sup>1</sup>
- So, we can always get at least one bit of information per intervention  $\Rightarrow$  a tree on p nodes takes at most  $\lceil \log_2 p \rceil$  interventions to orient

Theorem (Greenewald et al., 2019)

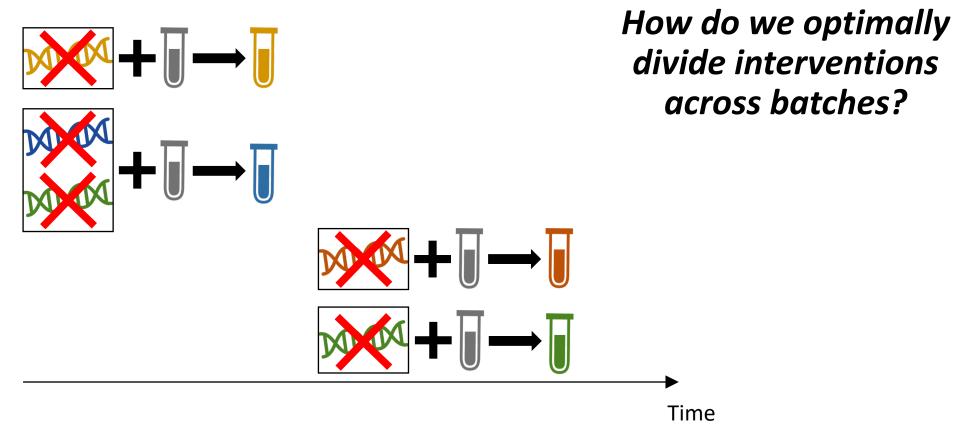
The expected number of interventions required by the central node algorithm is at most twice the expected number of interventions required by the optimal algorithm.

# Part IV: Targeted experimental design and other open challenges

#### Targeted experimental design

- So far, we've treated all structural causal information about our graph as equally important
- Rarely true!
  - We may only care about a subset of structural information (e.g., existence of certain paths)
    - <u>ABCD-Strategy: Budgeted Experimental Design for Targeted Causal Structure Discovery</u> (Agrawal et al., 2019)
  - We may only care about finding an "optimal" information for some goal, e.g. so that the interventional distribution is close to some target distribution
    - Matching a Desired Causal State via Shift Interventions
- For different learning goals, how does interventional and sample complexity change?

#### Batched experimental design



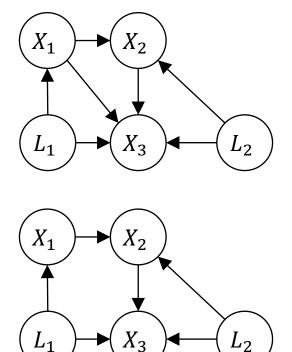
ABCD-Strategy: Budgeted Experimental Design for Targeted Causal Structure Discovery (Agrawal et al., 2019)

Batched bandit problems (Perchet et al., 2016)

#### Noisy interventional data

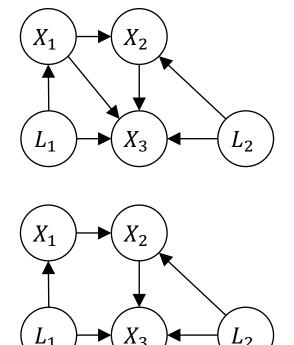
- How much better can we do than the "repeat until desired confidence" strategy for dealing with noise?
  - <u>Sample Efficient Active Learning of Causal Trees</u> (Greenewald et al., 2019) use a multiplicative weights algorithm in the binary setting
  - <u>Learning and Testing Causal Models with Interventions</u> (Acharya et al., 2018) study "goodness-of-fit" testing and other problems in the discrete setting
    - Given a true distribution  $f_*$  and a hypothesized distribution  $\tilde{f}$  both Markov to G, are all associated interventional distributions ( $f_*^k$  and  $\tilde{f}^k$  for  $I_k$  any do-intervention) with  $\epsilon$  total variation distance?

We can't always identify causal graphs in insufficient systems from just single node interventions



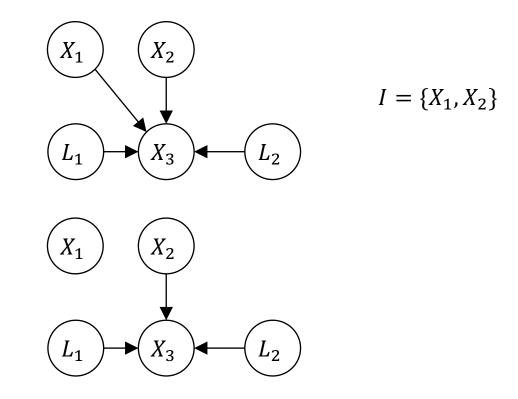
<sup>1</sup>Causation and Intervention (Eberhardt, 2008)

But we can identify them when the number of intervened nodes is unbounded



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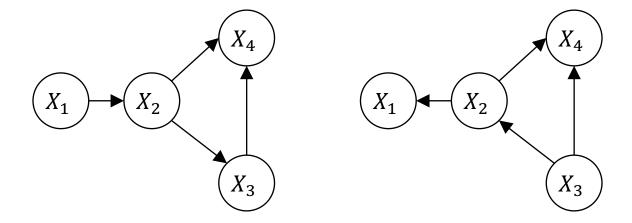
What graphs can we identify when the number of intervention targets is bounded by K?

#### Lower bounds on interventional complexity

- Suppose a friend tells you the true graph  $G^*$ , but you're skeptical and want to check it, while using the minimal number of single-node interventions
- If  $G^*$  is a tree: only 1 intervention is required (at the root)
- If  $G^*$  is a clique on p nodes:  $\lfloor \frac{p}{2} \rfloor$  interventions are required
- Active Structure Learning of Causal DAGs via Directed Clique Trees (Squires et al., 2020) give a polynomial-time algorithm for finding the "minimal verifying intervention set" (MVIS) for any DAG  $G^*$ , the size of which is denoted  $m(G^*)$

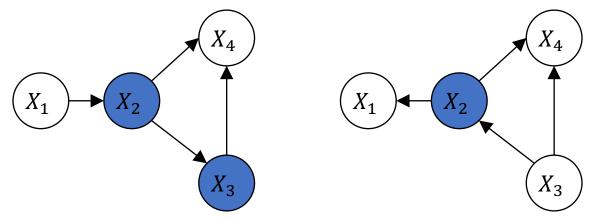
#### Lower bounds on interventional complexity

•  $m(G^*)$  is not necessarily the same across a Markov equivalence class



#### Lower bounds on interventional complexity

•  $m(G^*)$  is not necessarily the same across a Markov equivalence class



• Squires et al. (2020) give a "universal" lower bound on  $m(G^*)$  that holds across the equivalence class, improved in Almost Optimal Universal Lower Bound for Learning Causal DAGs with Atomic Interventions (Porwal et al., 2021)