

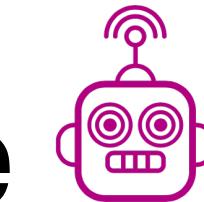
# General Identifiability with Arbitrary Surrogate Experiments

**Sanghack Lee**  
with Juan Correa and Elias Bareinboim

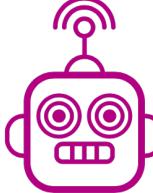
Columbia University

**AAAI 2020**  
(presented at UAI 2019)

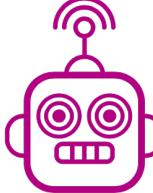
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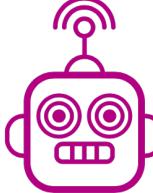
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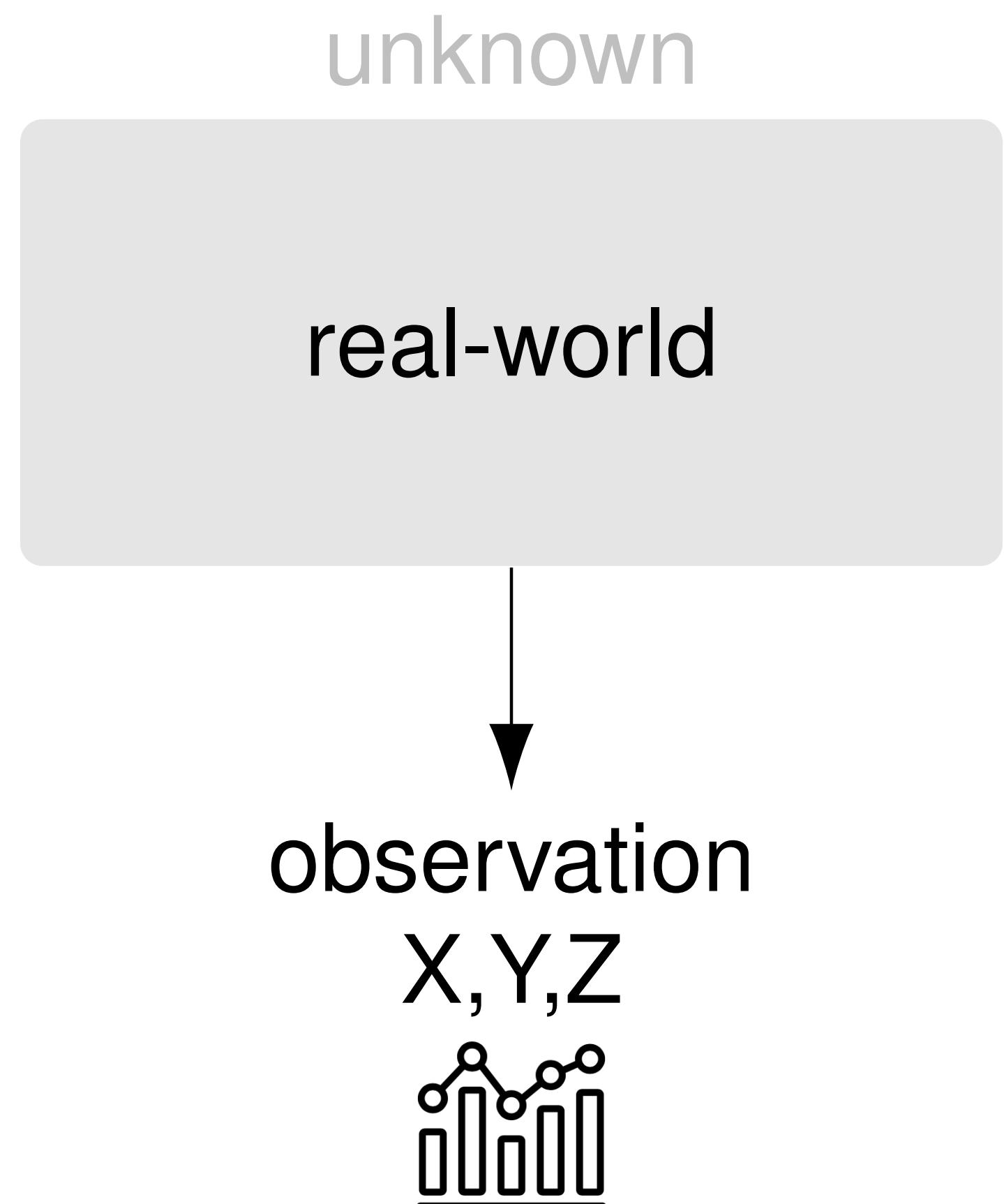
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- Causal Effect **Identifiability** concerns about precisely determining the effect of intervention given information (e.g., causal assumptions and an observational data).
- General **Identifiability** considers identifying a causal effect given an arbitrary combination of observational and experimental data .
- We provided a graphical **necessary and sufficient condition** under which a causal effect of interest can be estimable. We devised a **sound and complete algorithm** which outputs a formula for the causal effect made with probabilities obtained from available data.

# Understanding Data

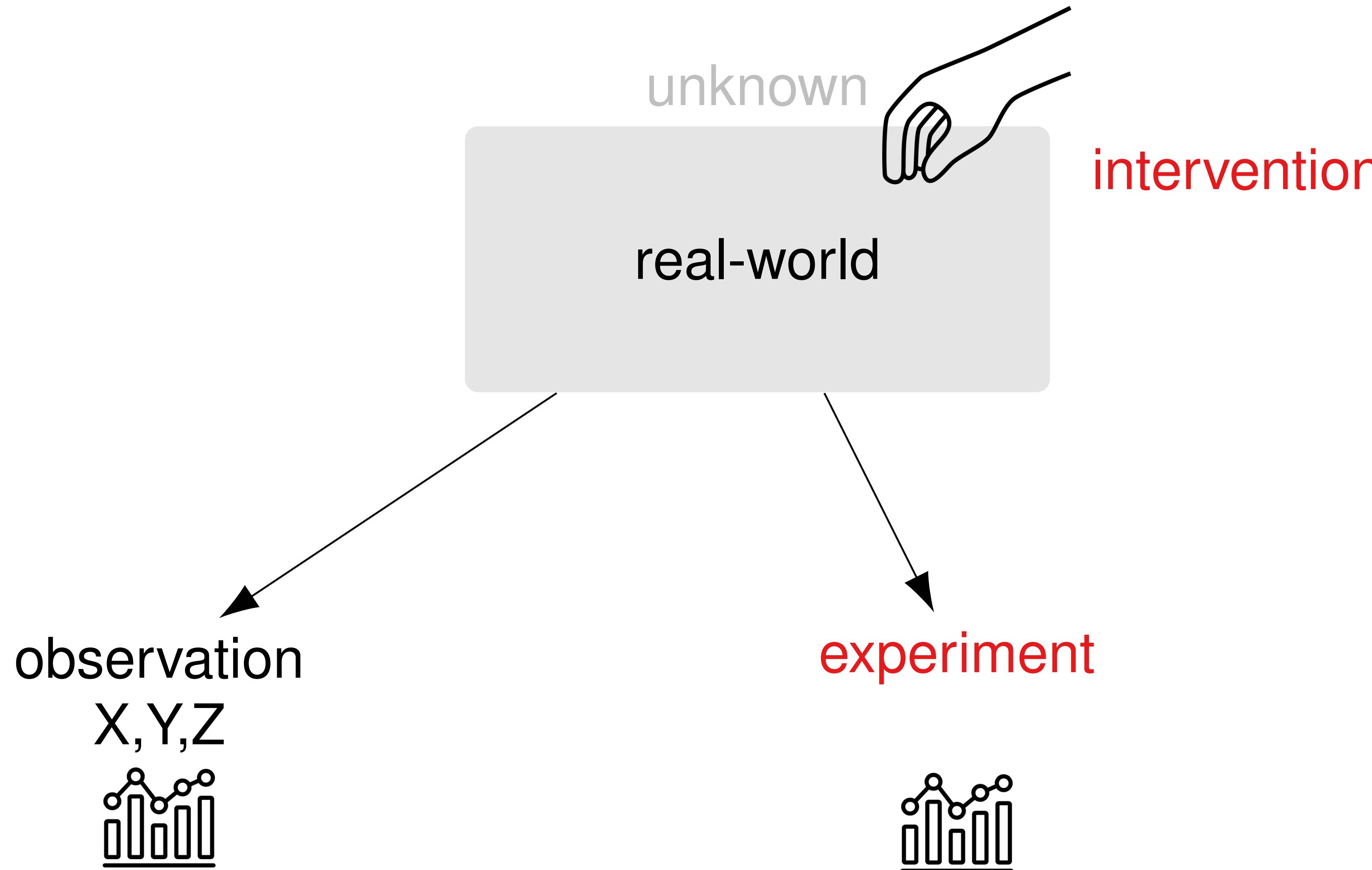
unknown

real-world

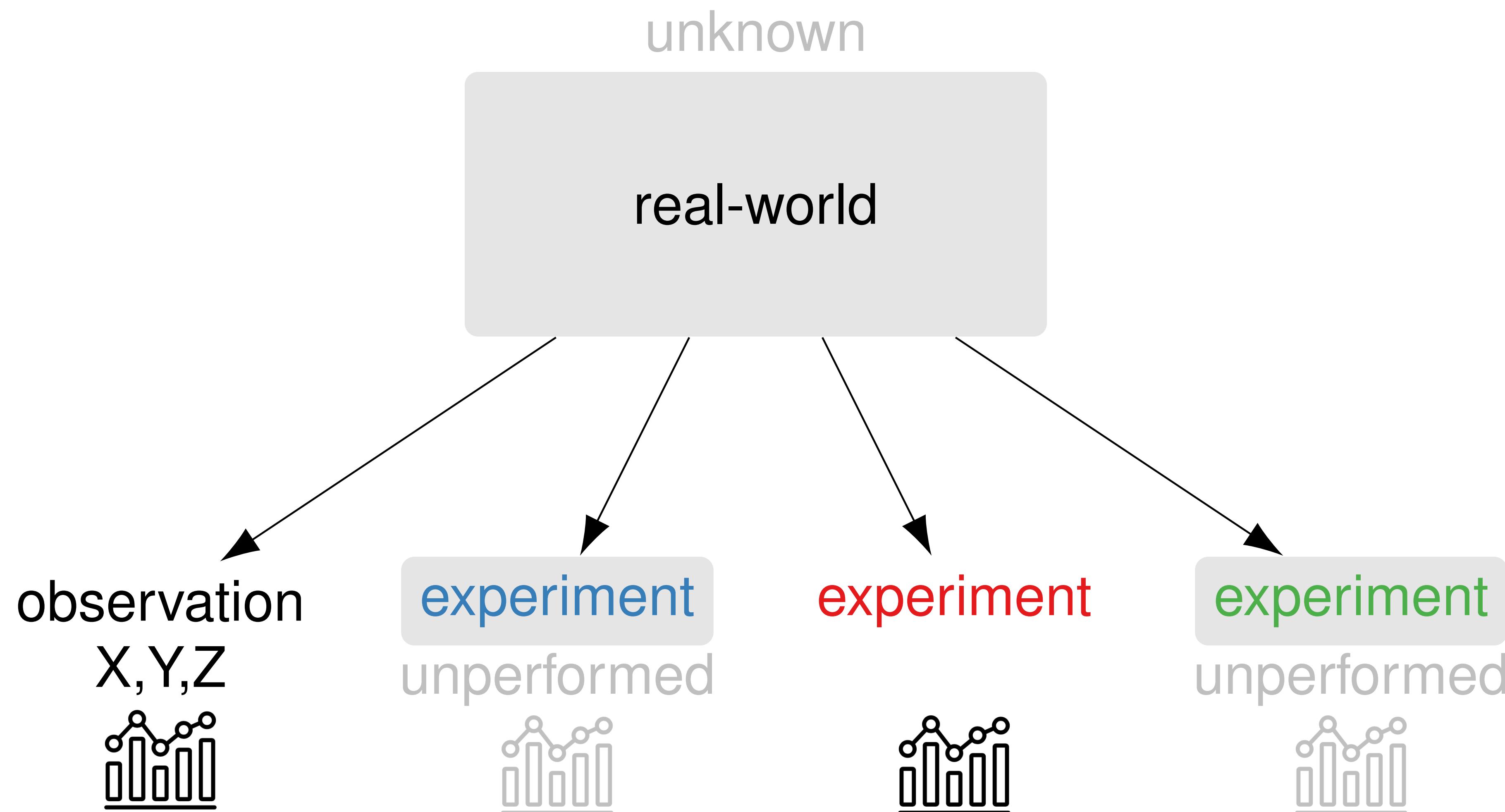
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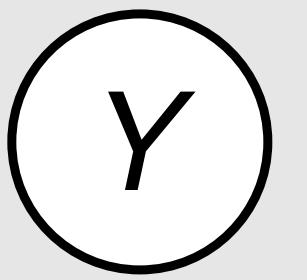
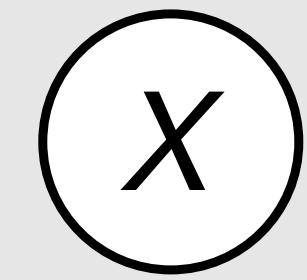


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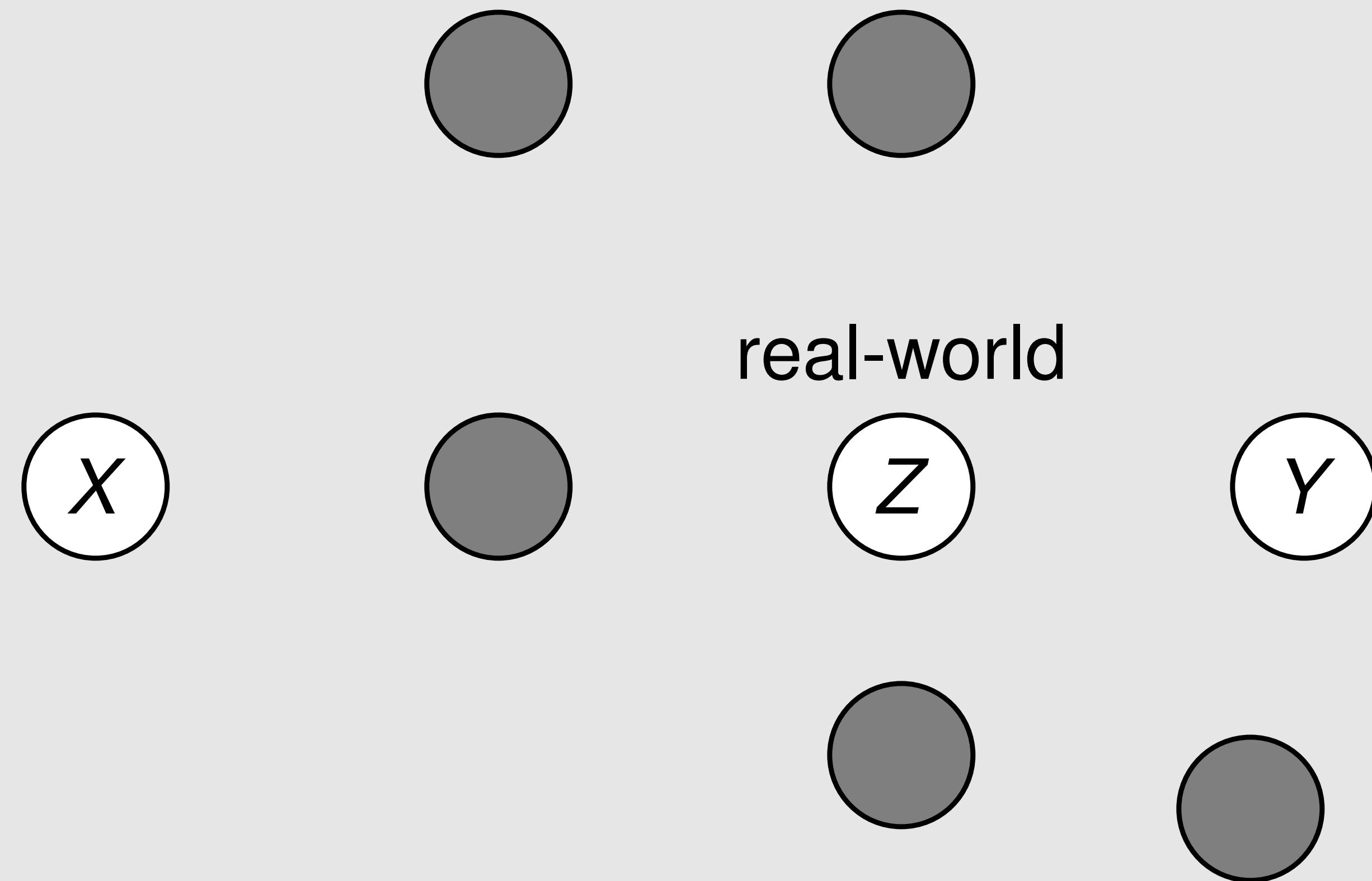


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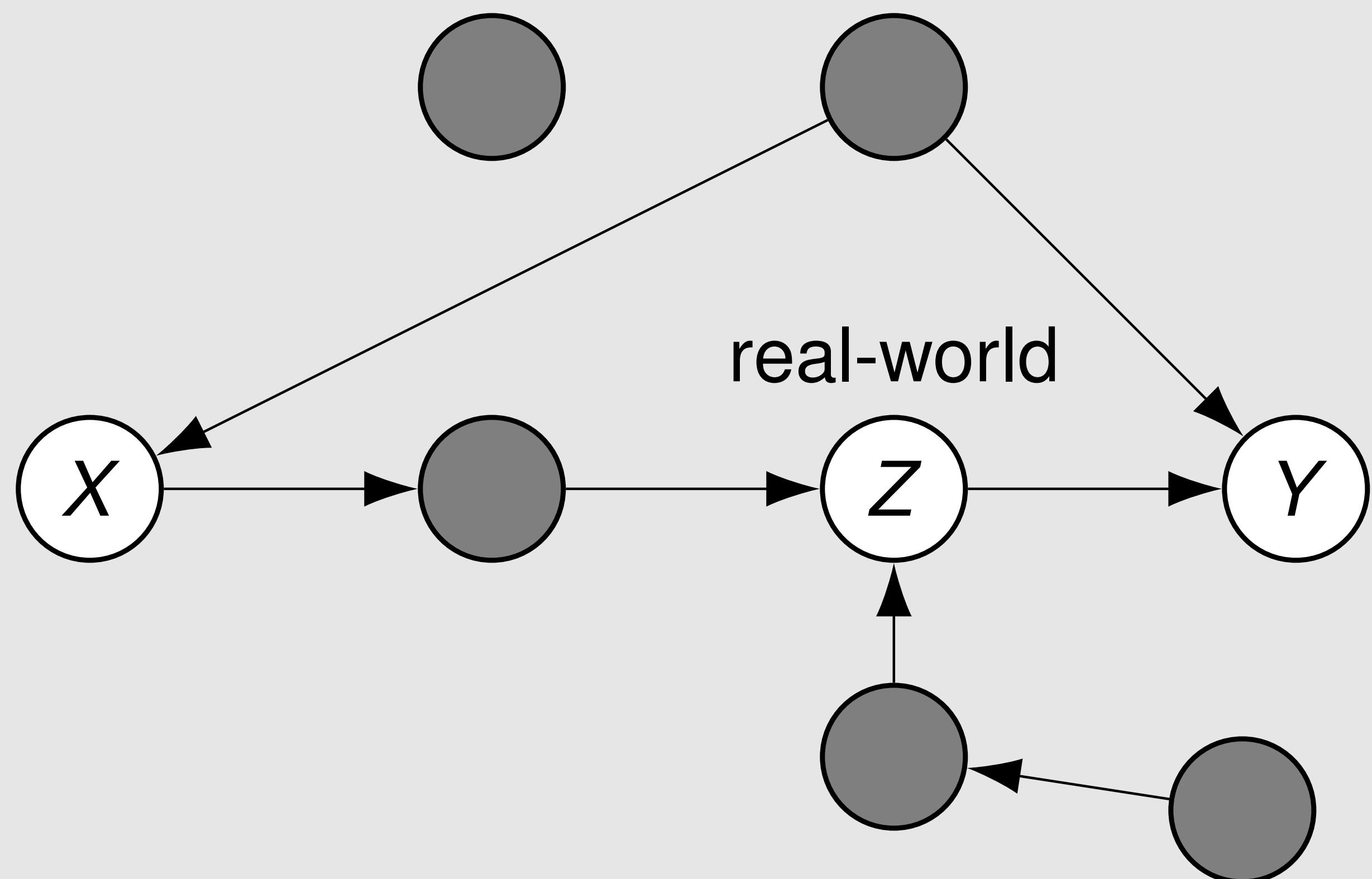
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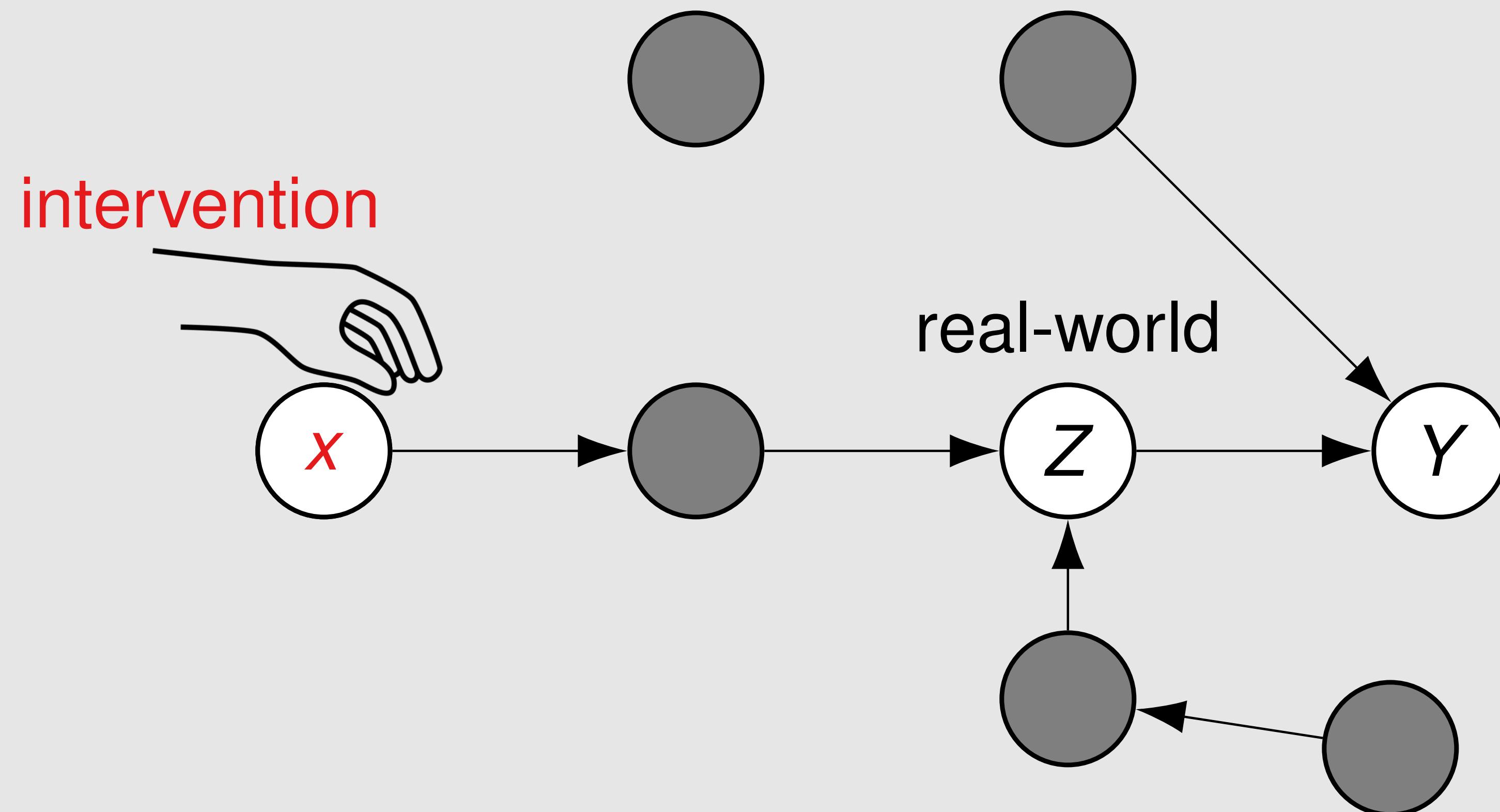
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# Causal Framework

SCM provides an abstraction of causality in the real-world.

## Definition (Structural Causal Model (Pearl))

SCM  $\mathcal{M}$  is a 4-tuple  $\langle \mathbf{U}, \mathbf{V}, \mathbf{F}, P(\mathbf{U}) \rangle$

- $\mathbf{U} = \{U_1, \dots, U_m\}$  are **exogenous** variables;
- $\mathbf{V} = \{V_1, \dots, V_n\}$  are **endogenous** variables;
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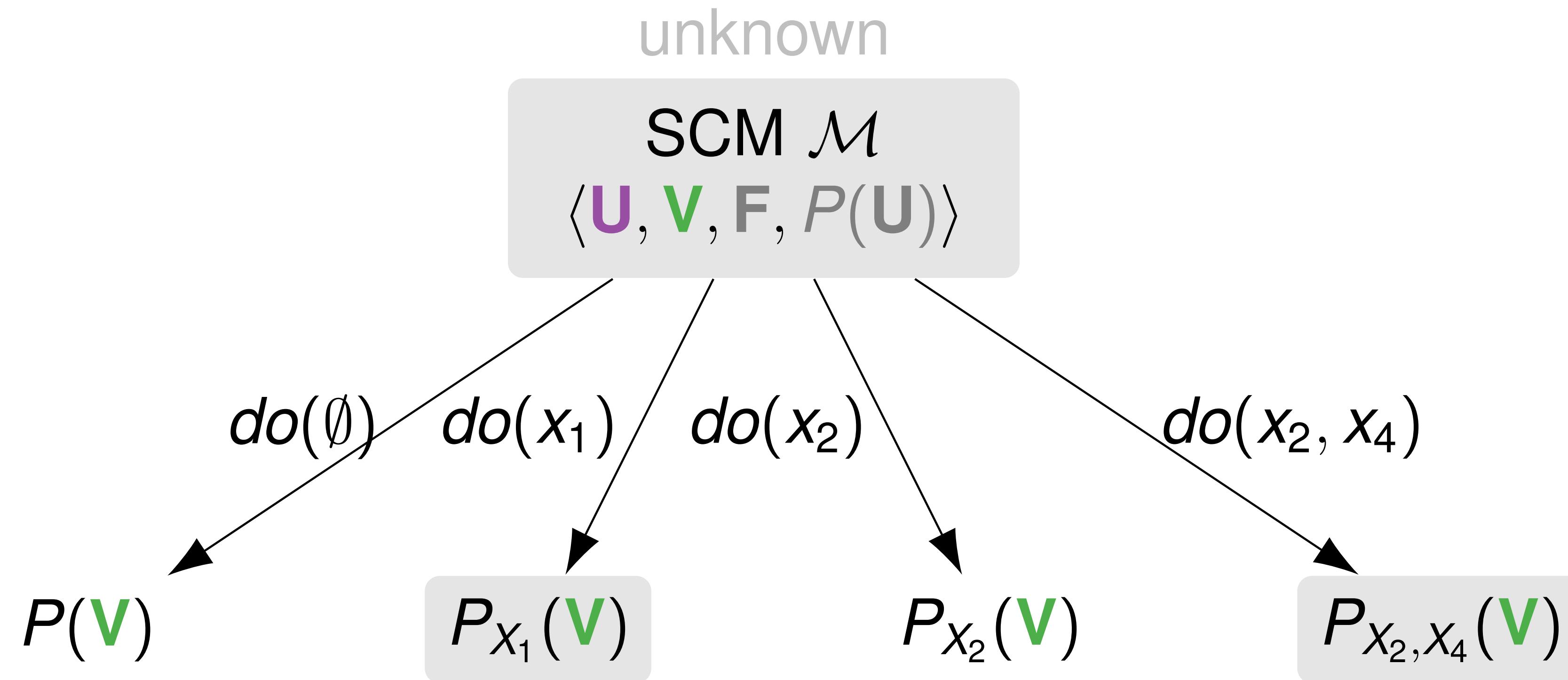
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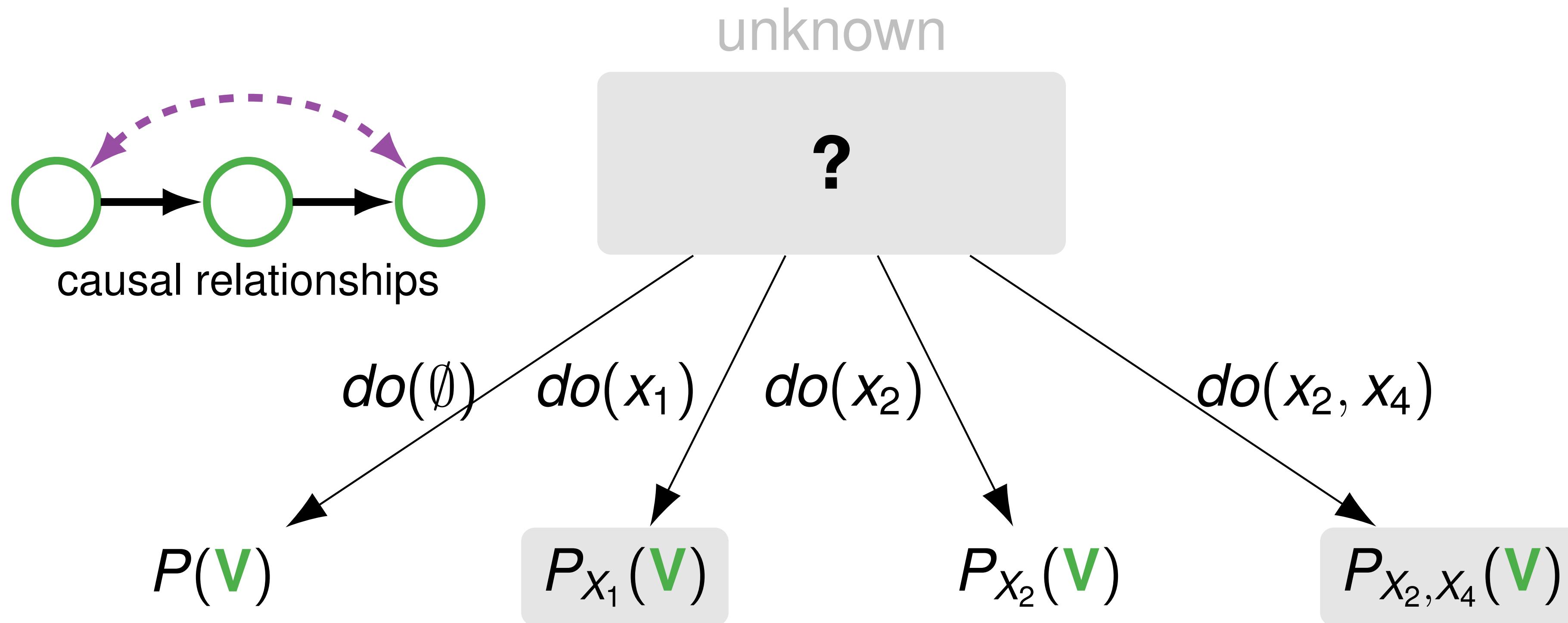
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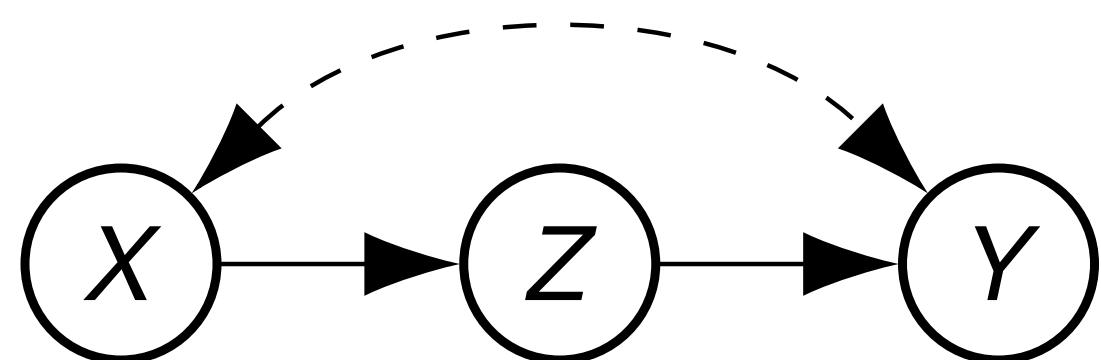


# (Classic) Causal Effect Identifiability

1 Query  $Q$

$$P_{\mathbf{x}}(\mathbf{y}) = P(\mathbf{y}|do(\mathbf{x}))$$

2 Causal Diagram  $\mathcal{G}$

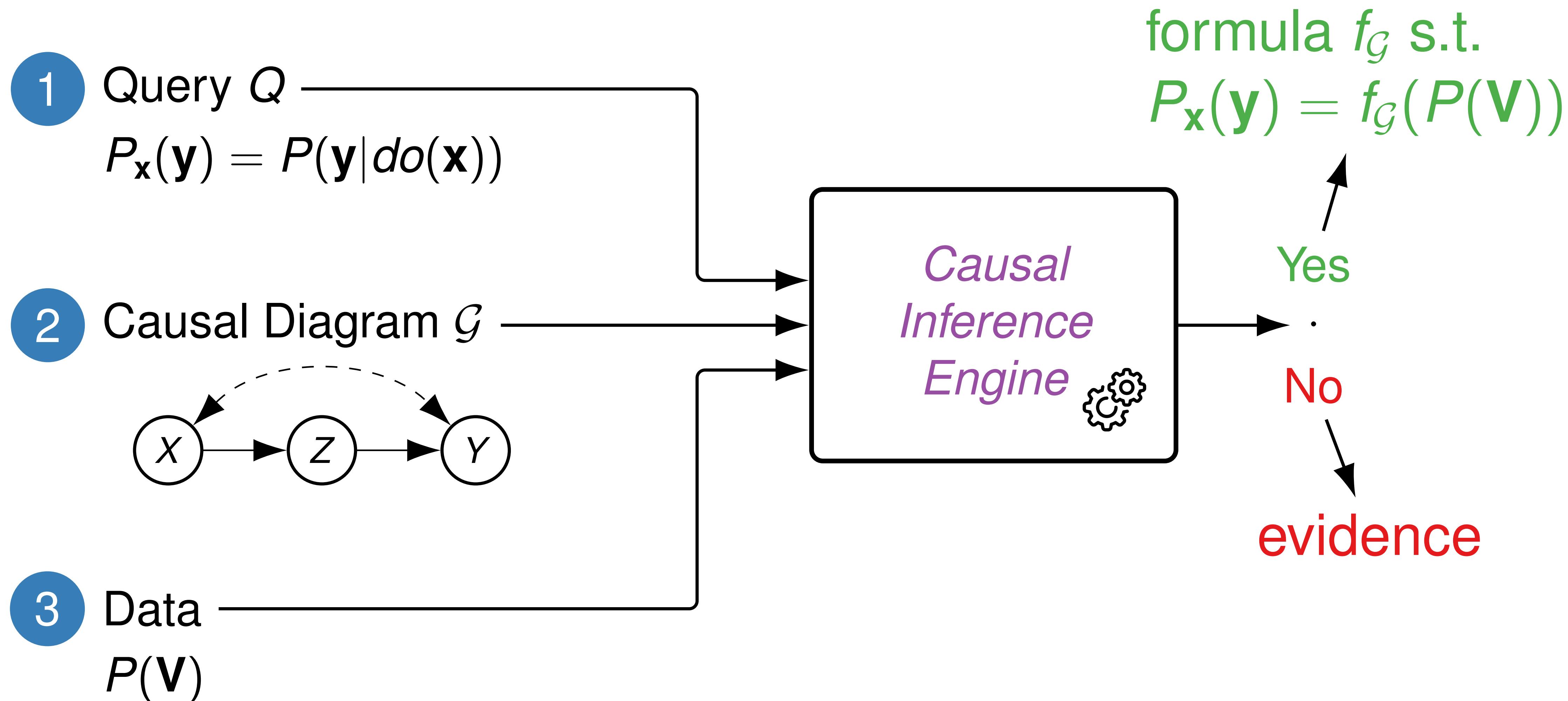


3 Data

$$P(\mathbf{V})$$

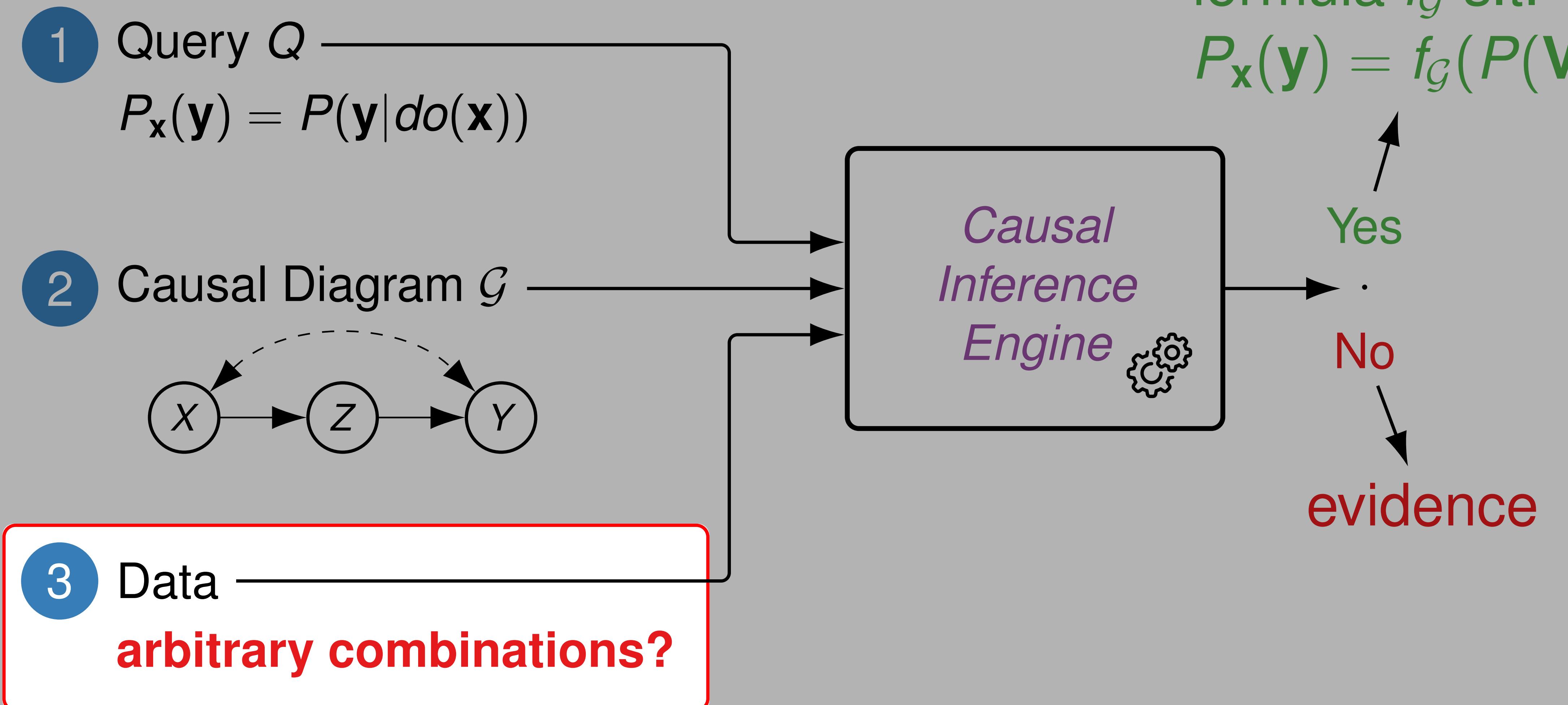
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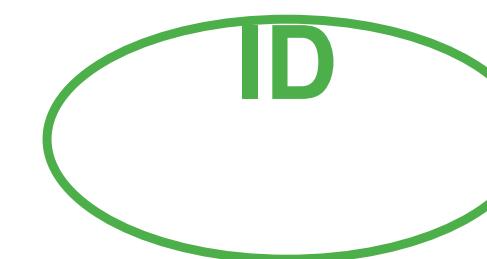
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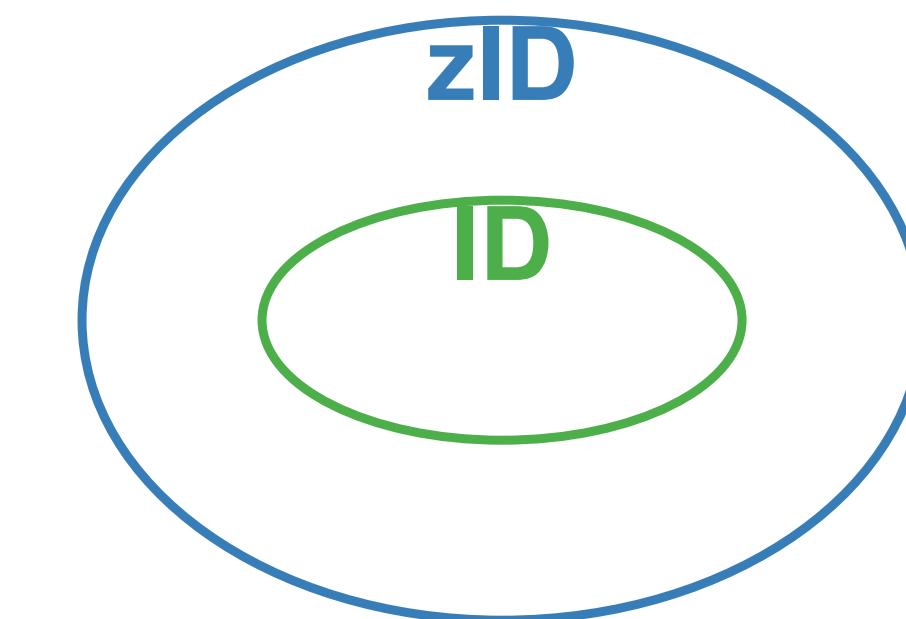
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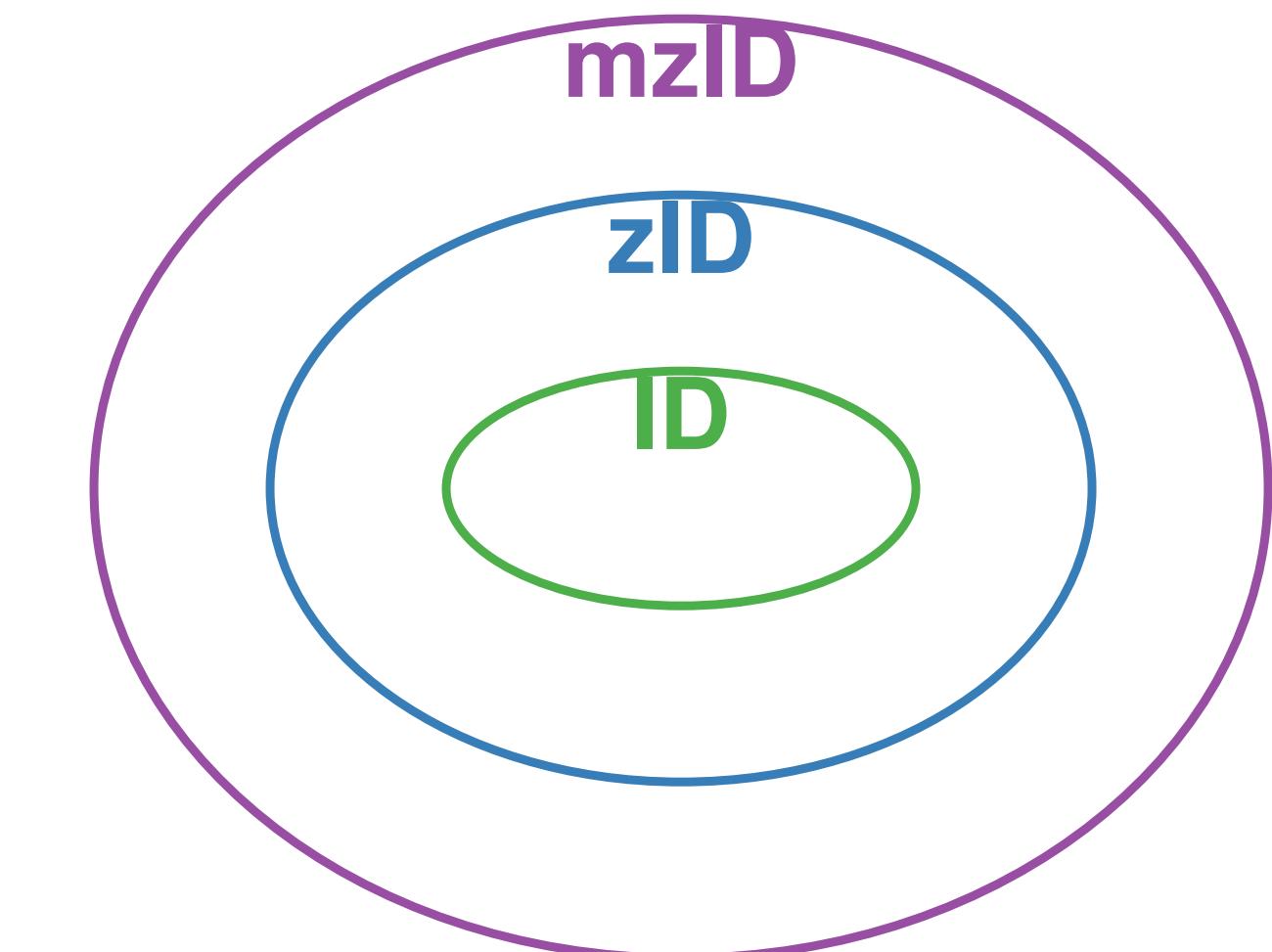
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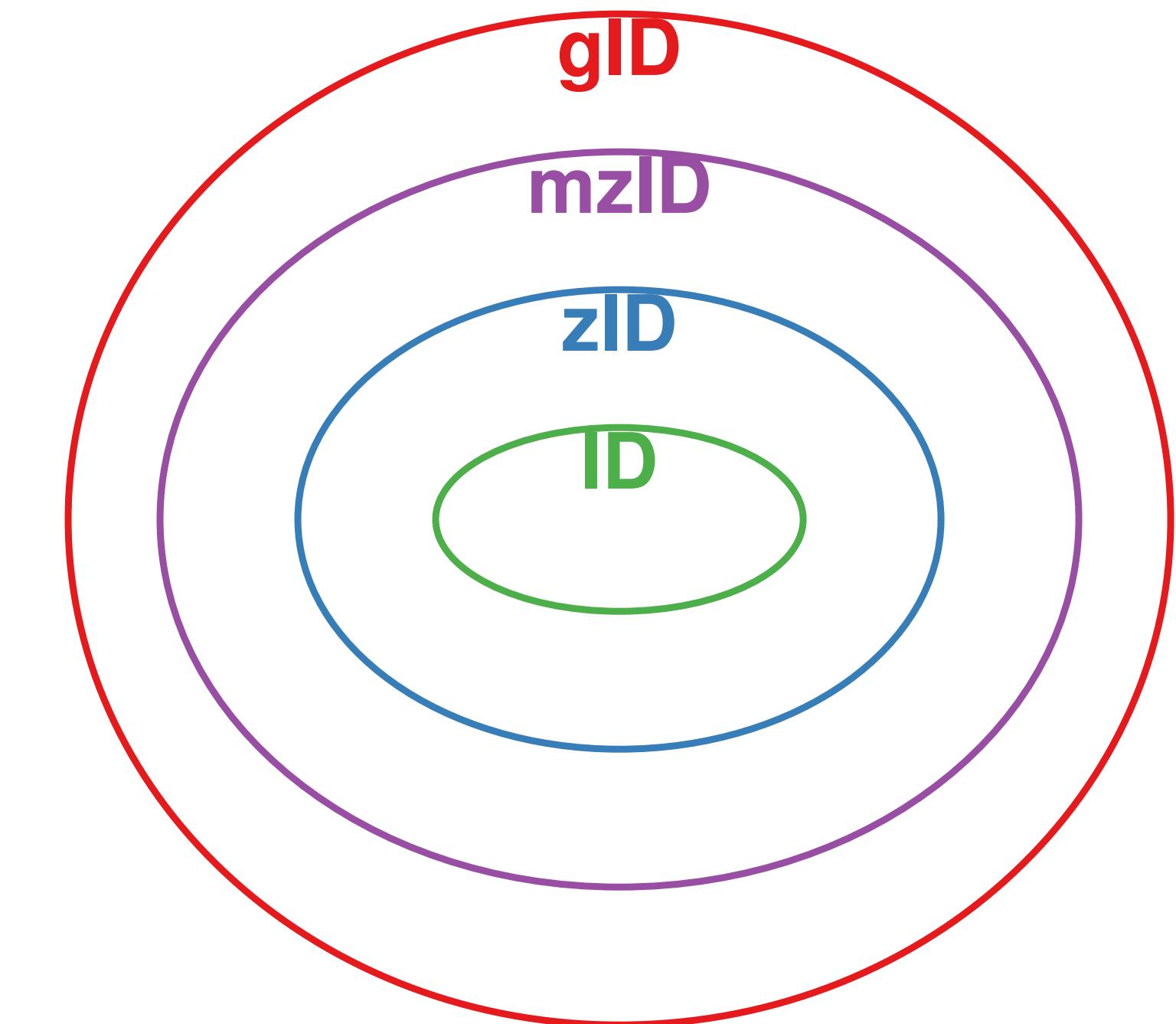
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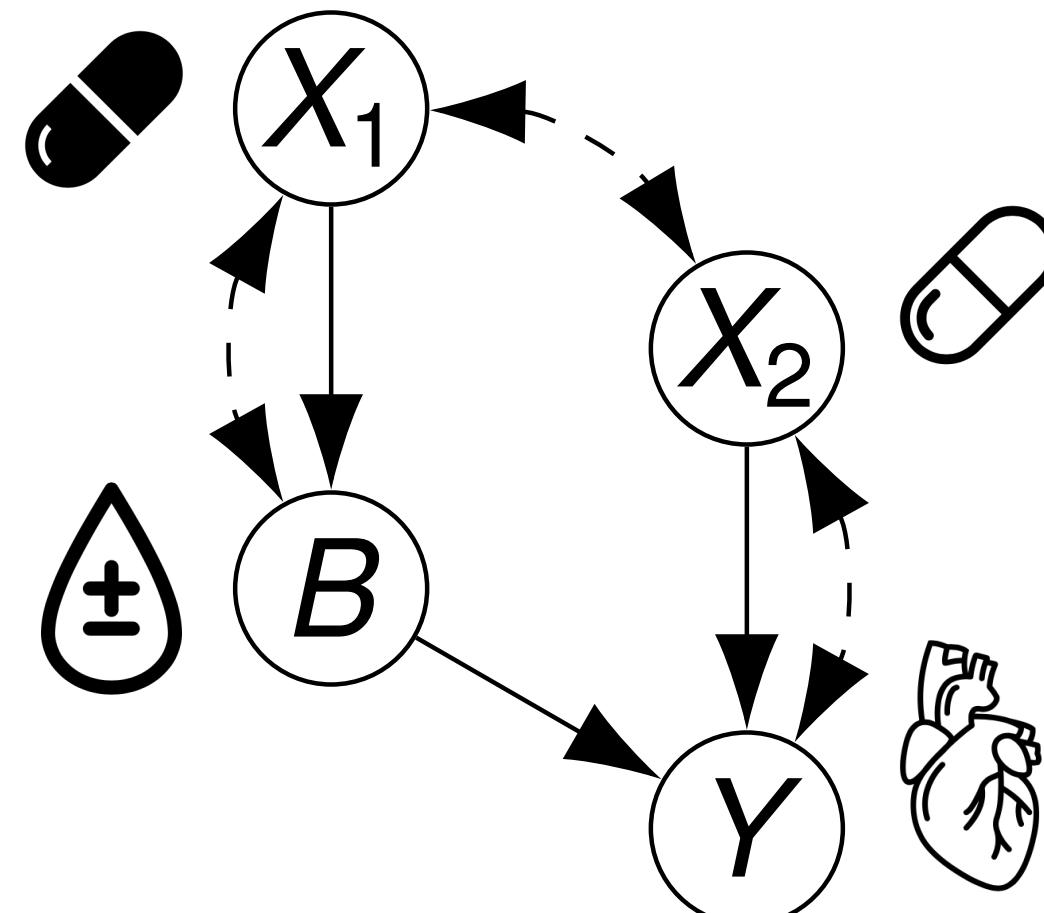
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**gID** A collection of arbitrary experiments [LCB'19]:

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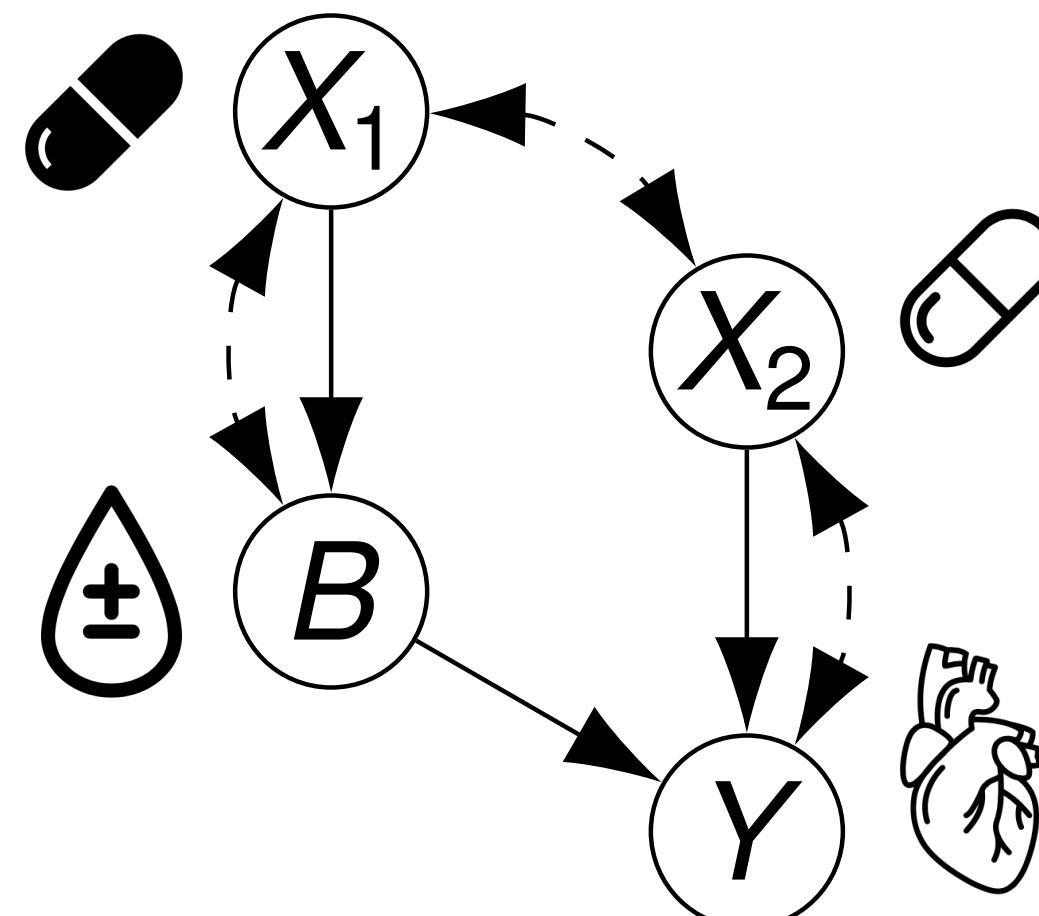
# Example: Drug-Drug Interactions



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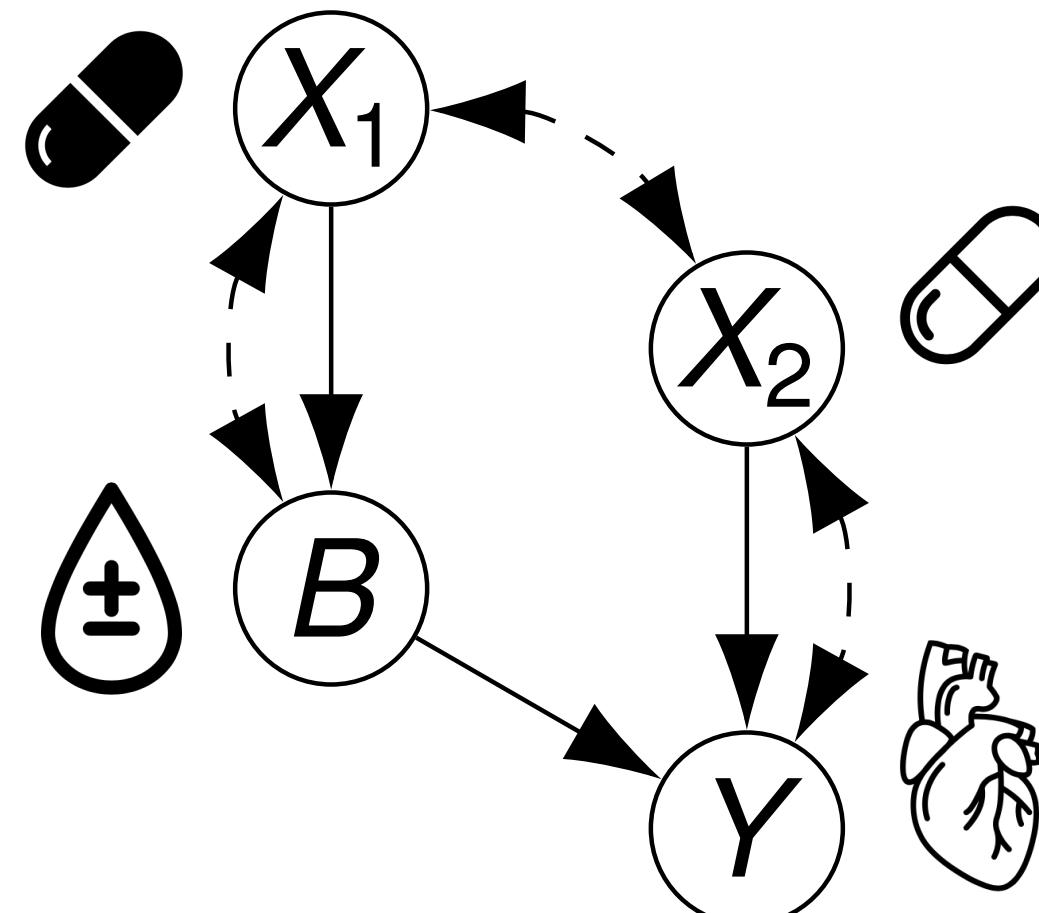
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$$P_{x_1, x_2}(y) \Leftarrow \{P_{x_1}(\mathbb{V}), P_{x_2}(\mathbb{V})\}?$$

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**Goal:** assess the effect of prescribing **both** treatments (capsule icons) on the risk of cardiovascular diseases from **individual** drug experiments, either capsule icon or circle icon.

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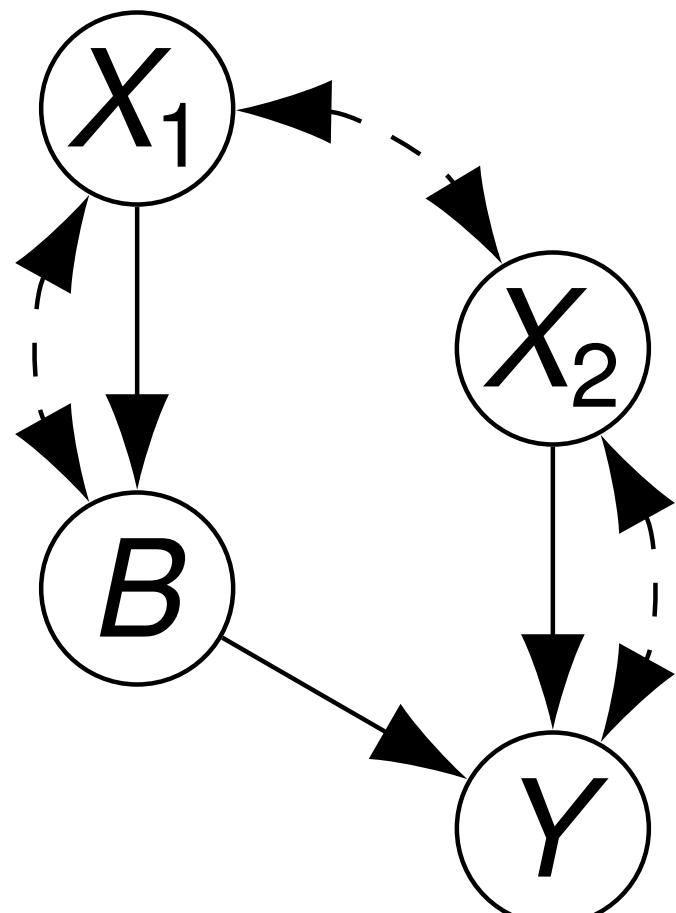
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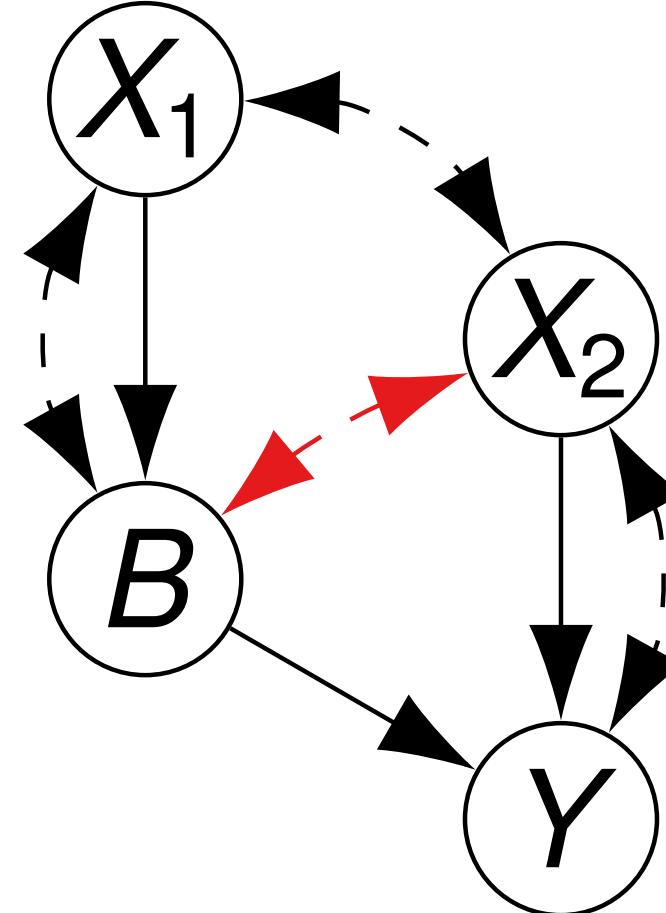
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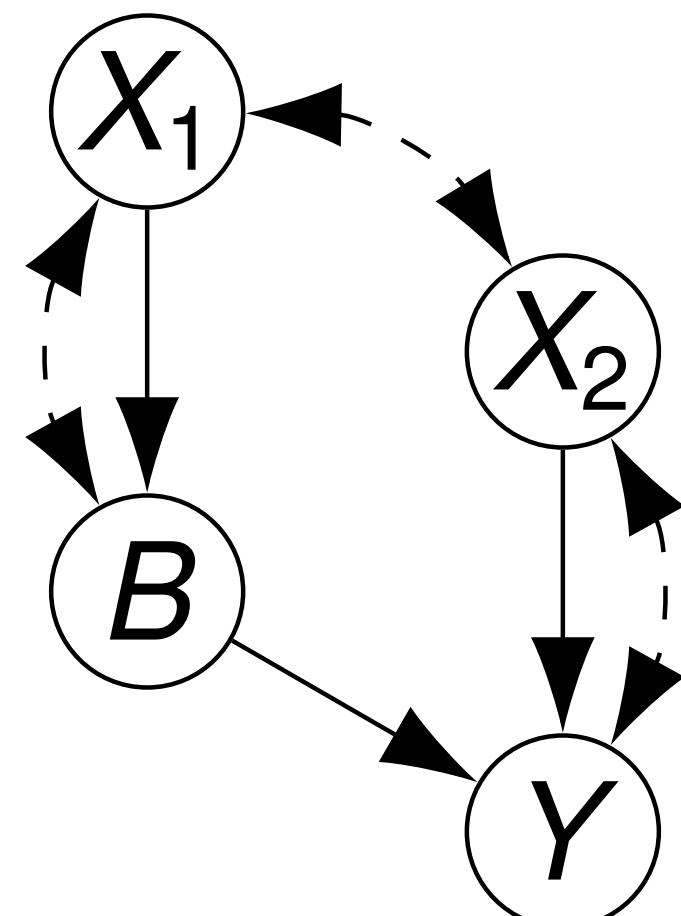


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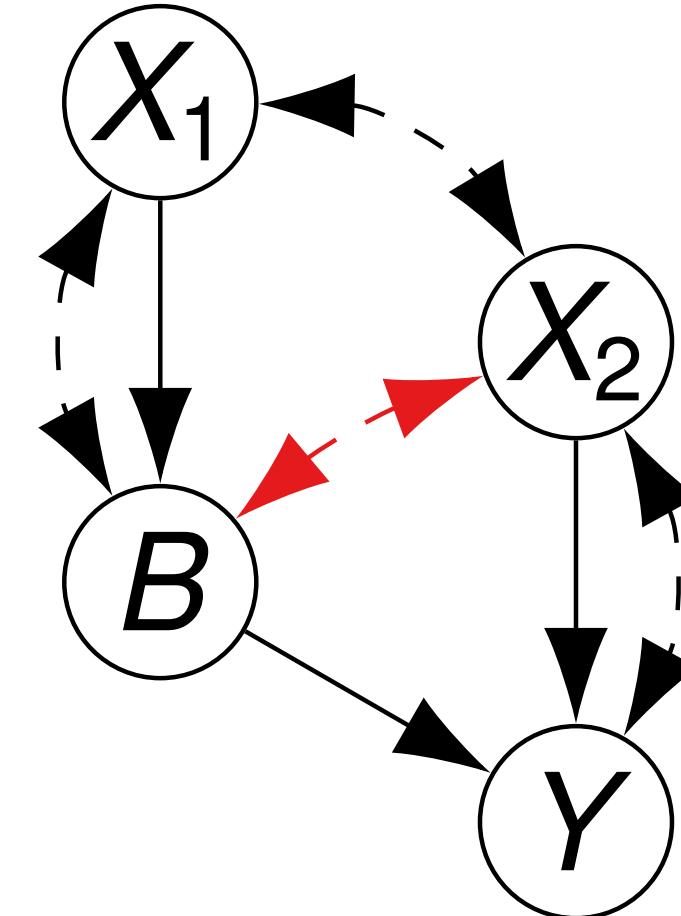
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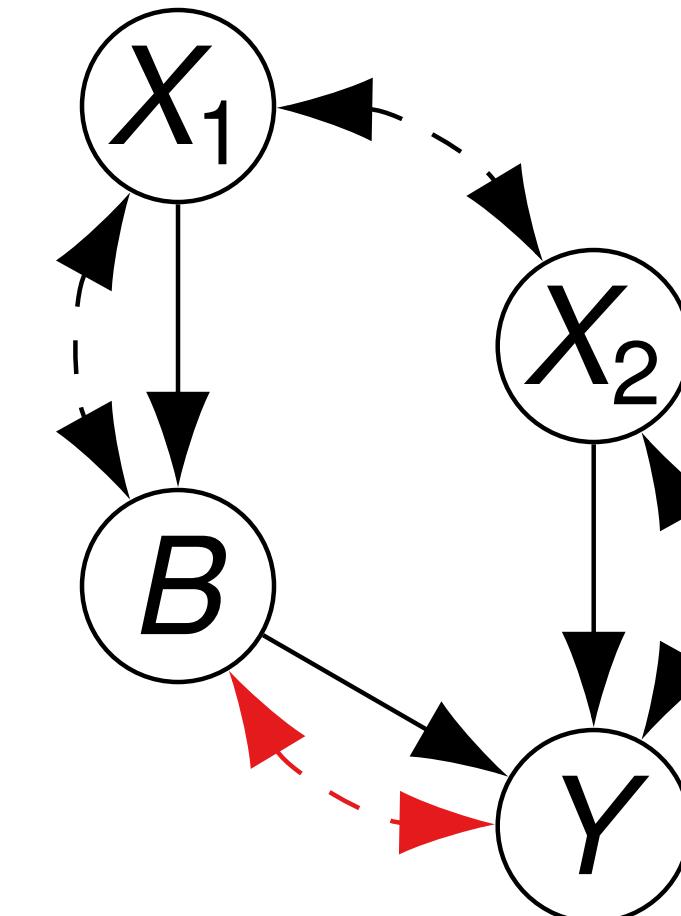
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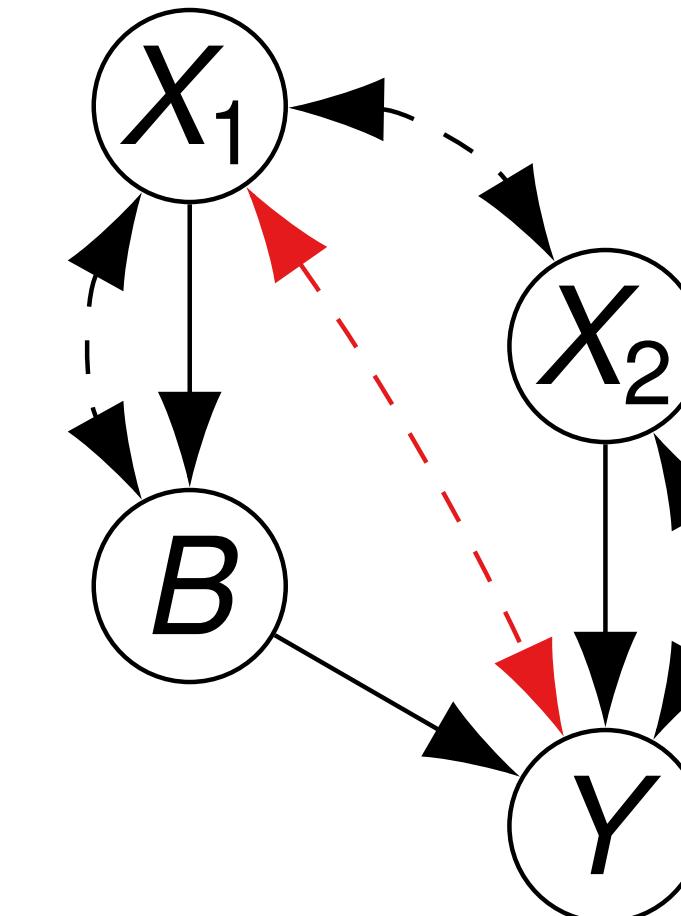
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(c) ✗



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# g-identifiability

- a sound algorithm

# Algorithm for gID

We developed a two-phase algorithm w/ probability axioms & do-calculus:

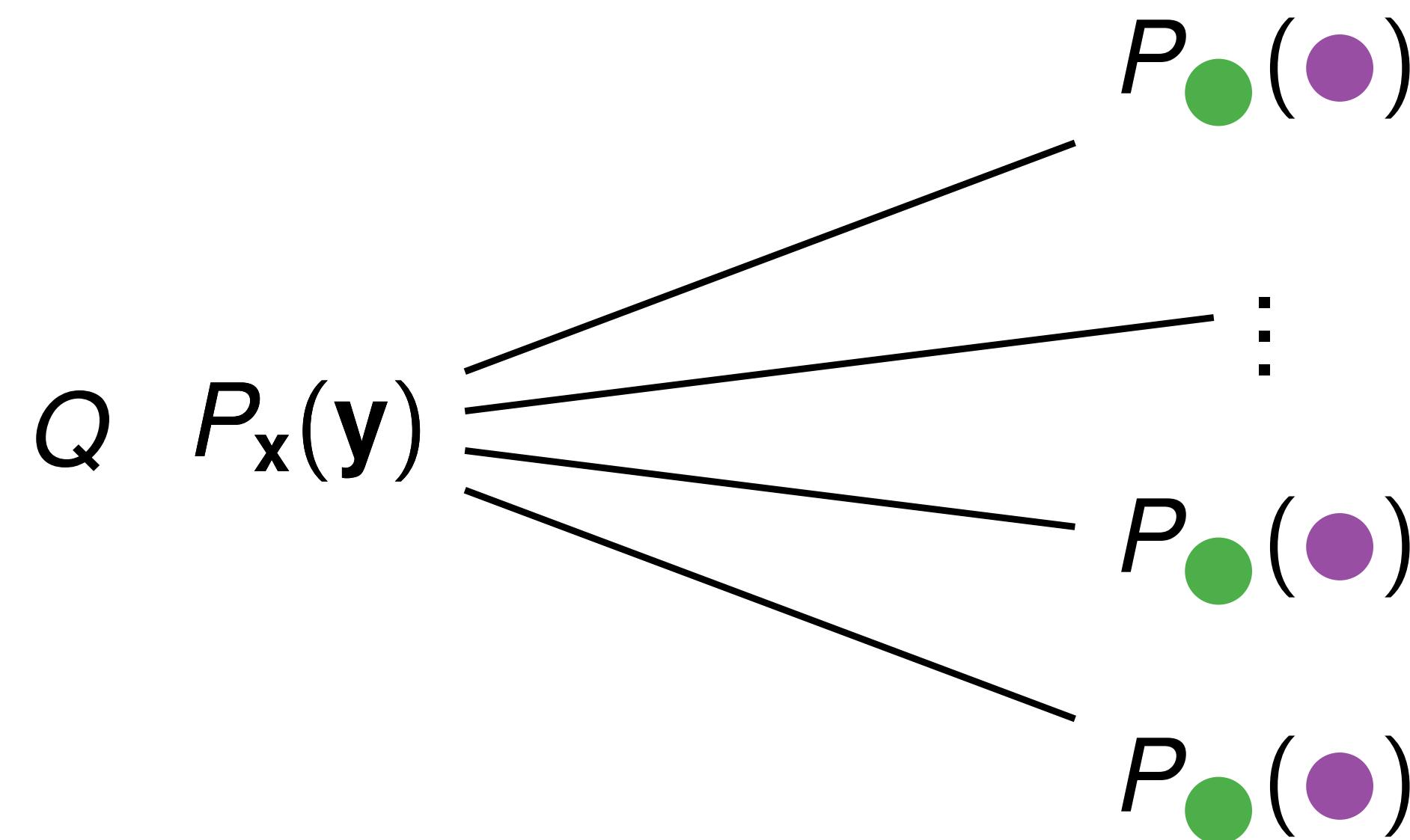
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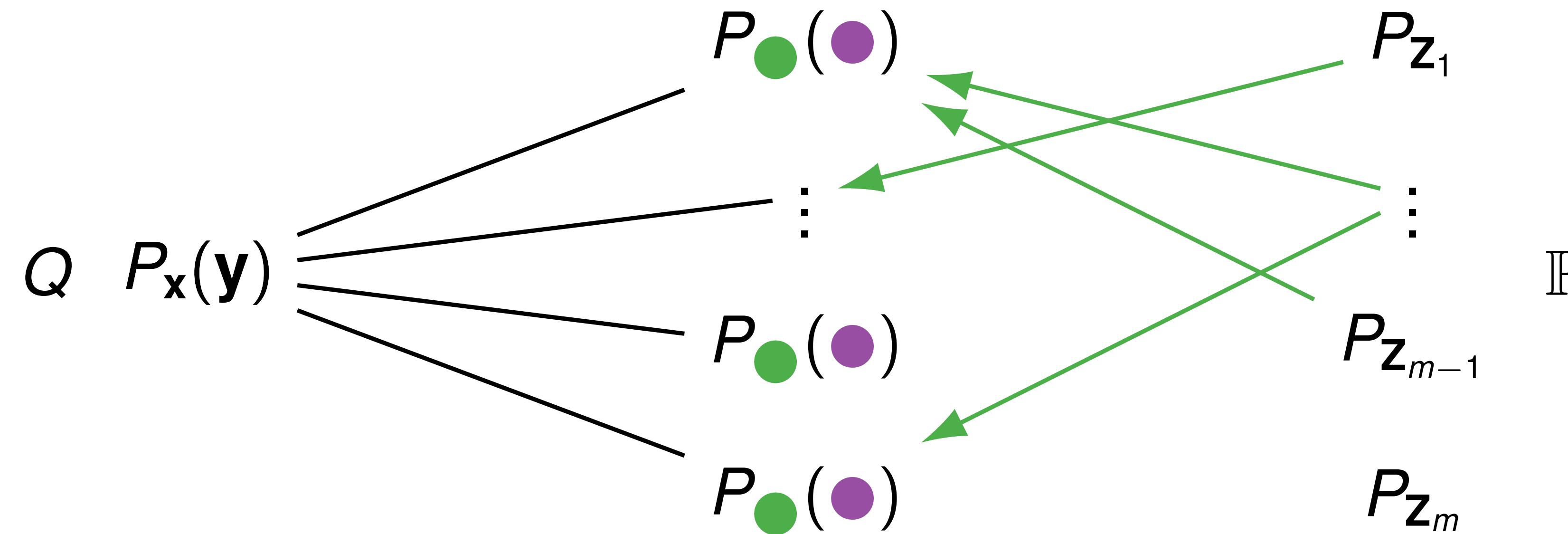


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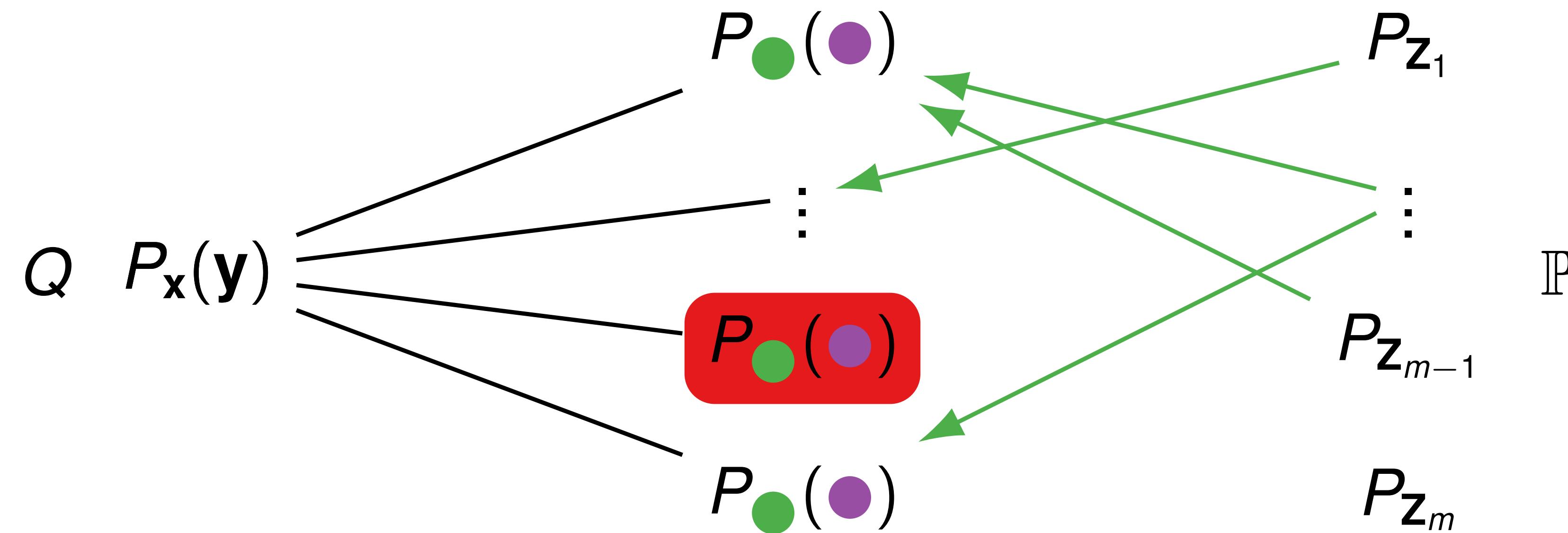
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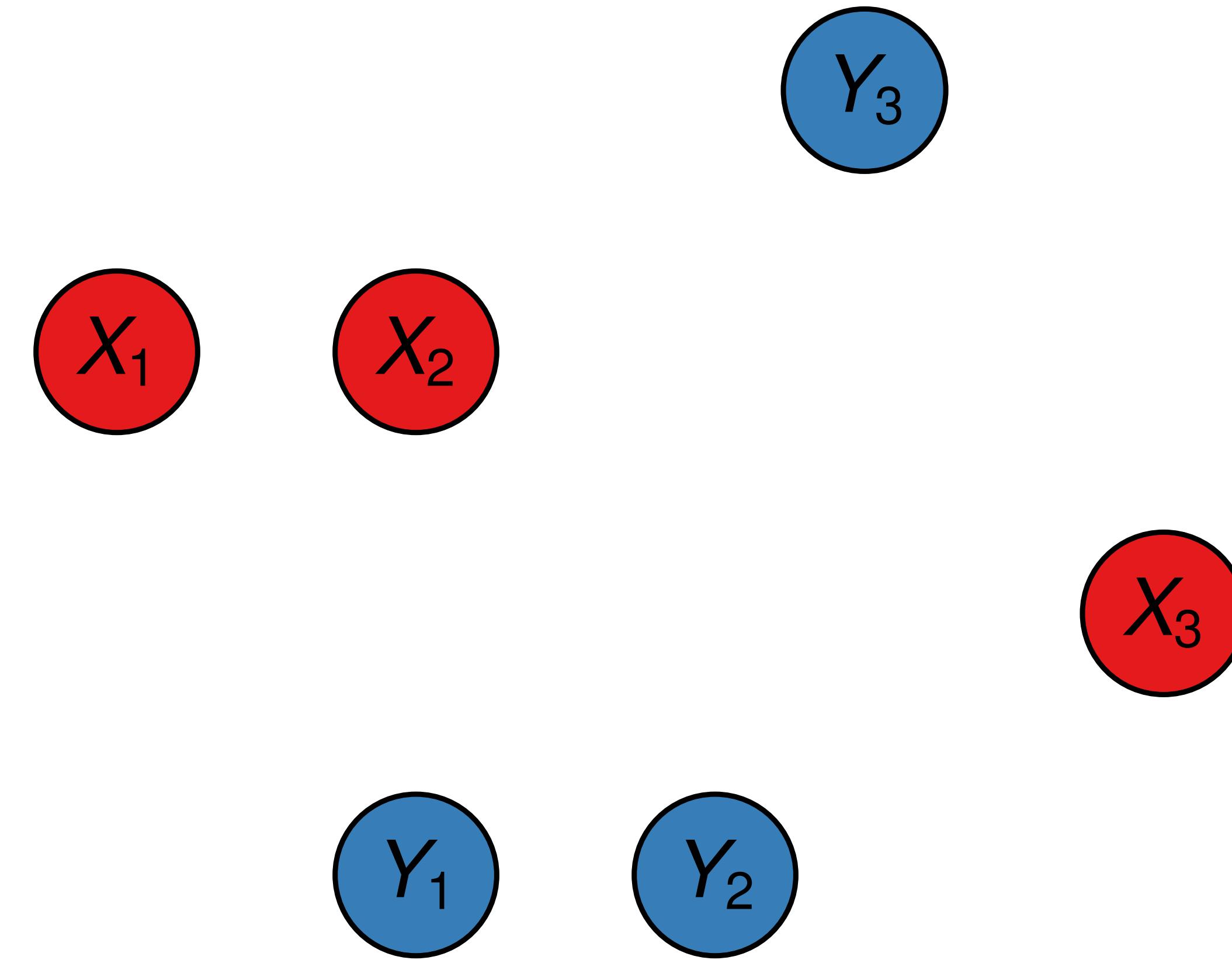
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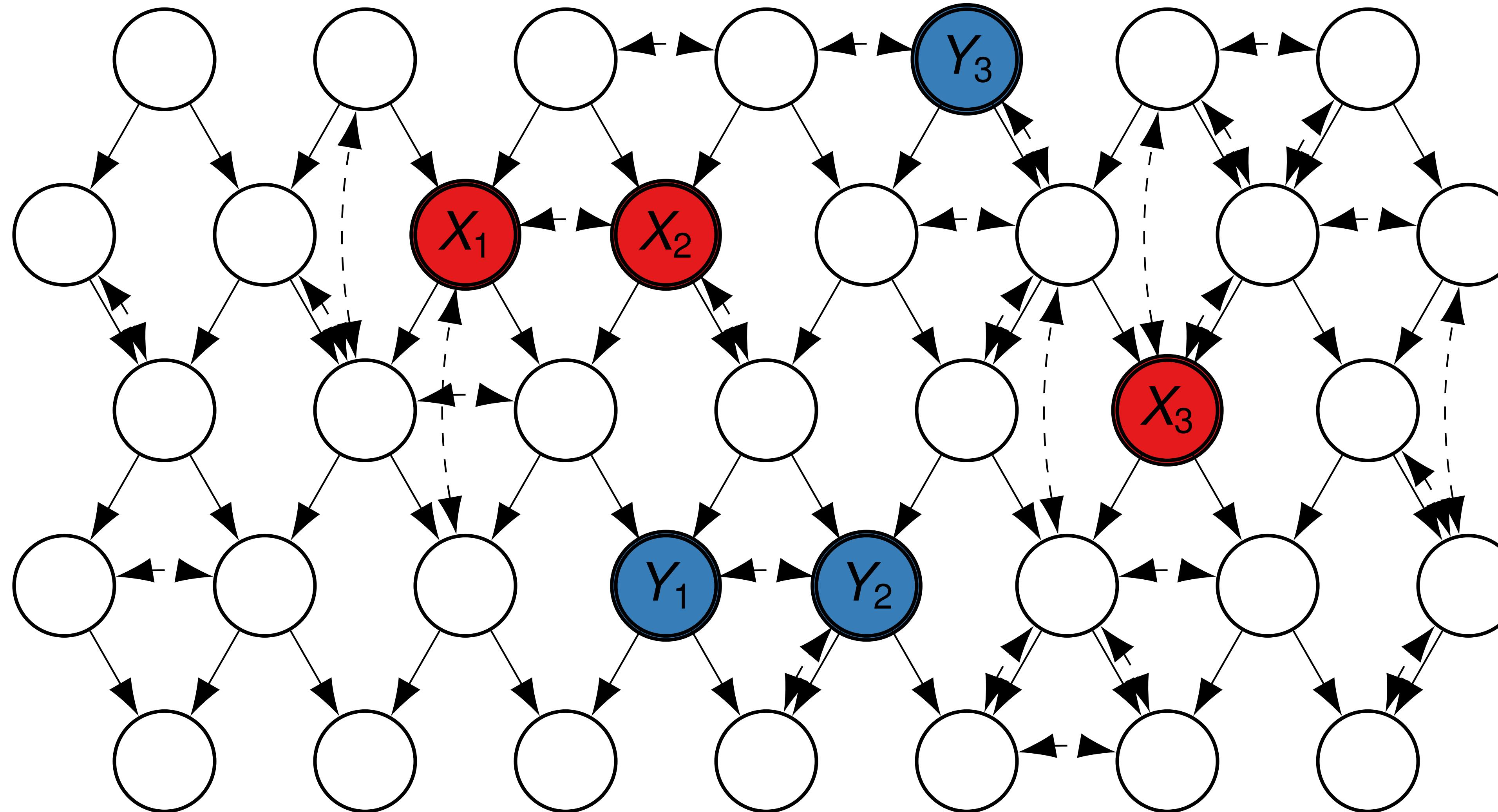
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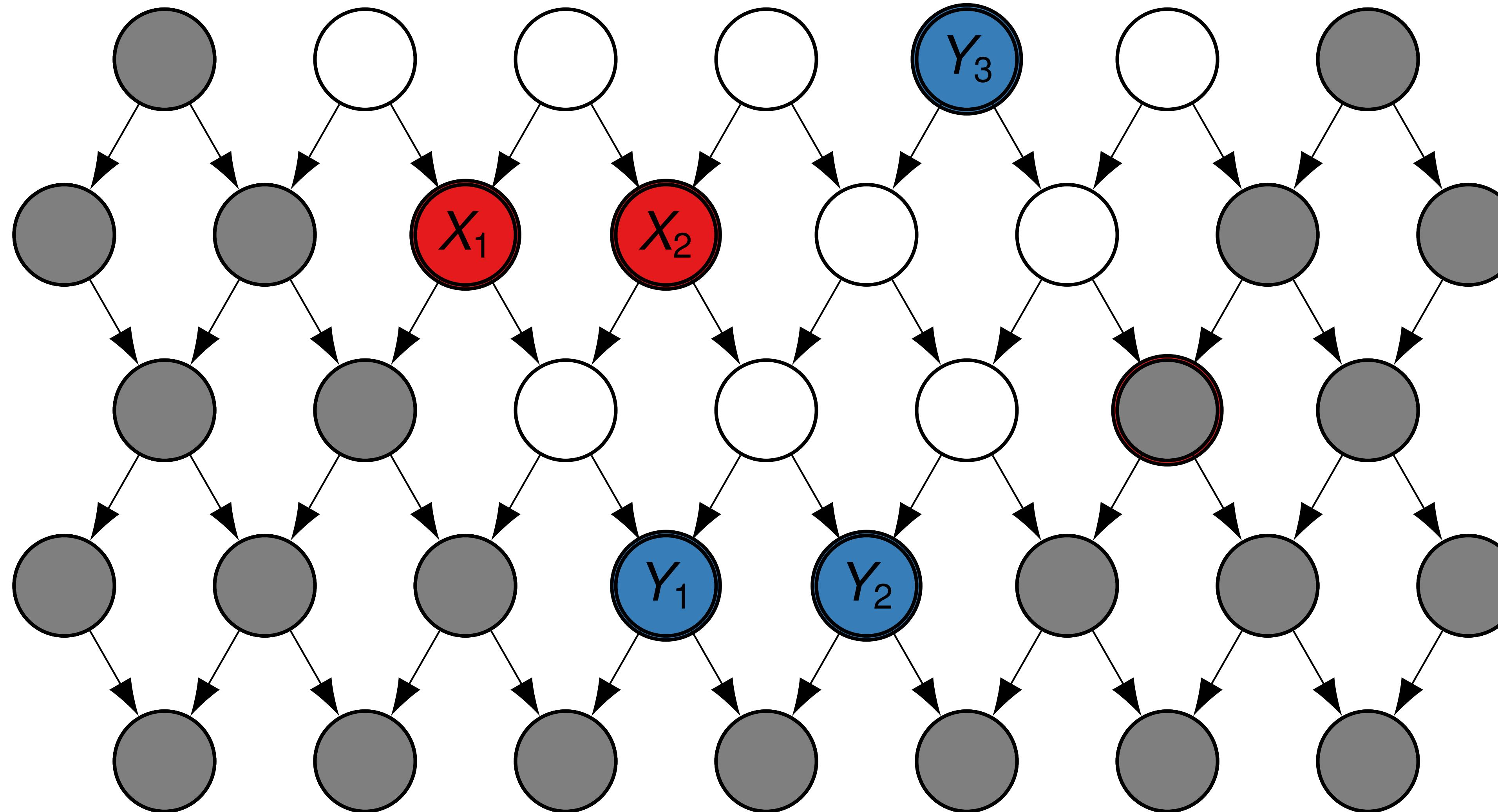
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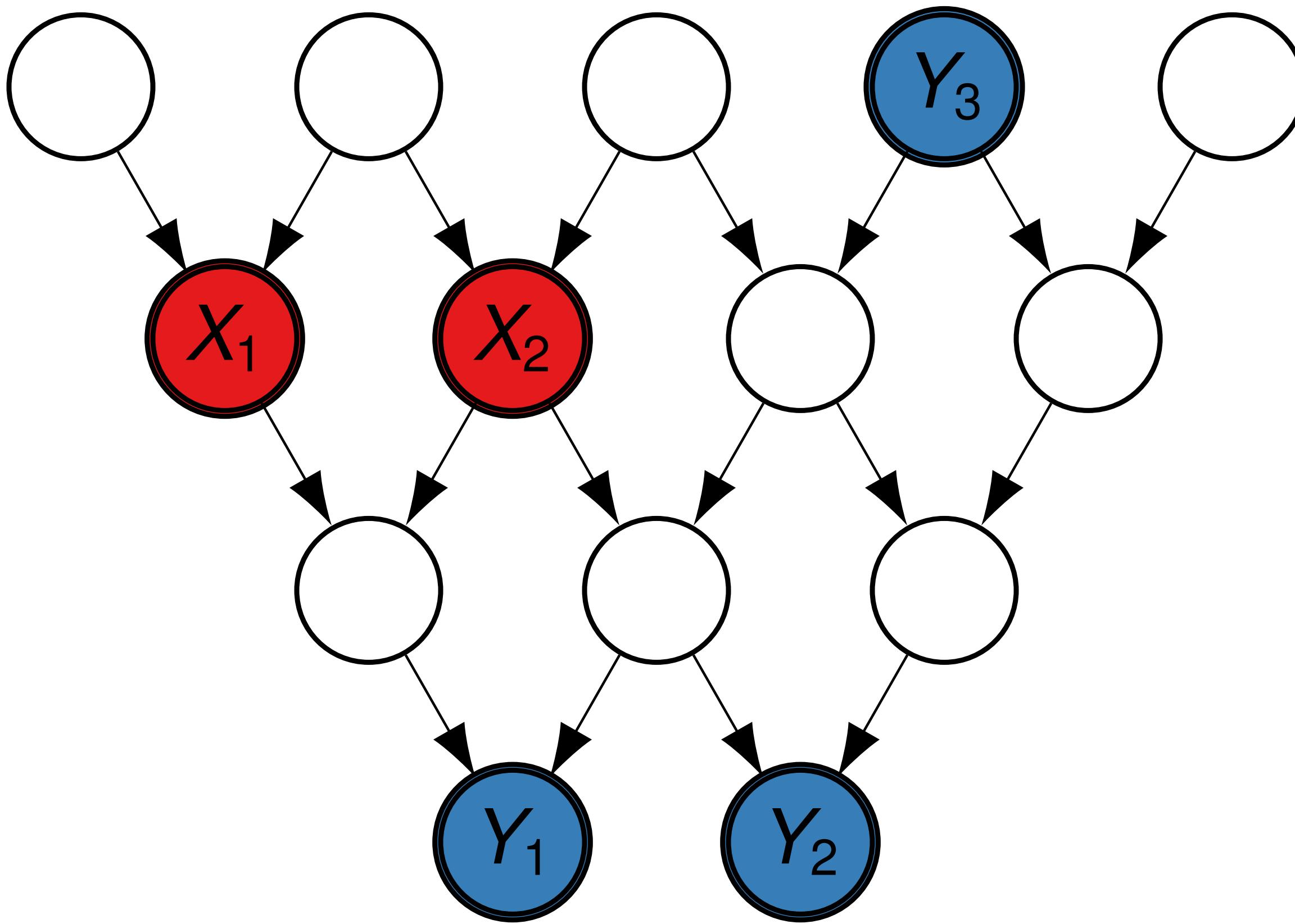
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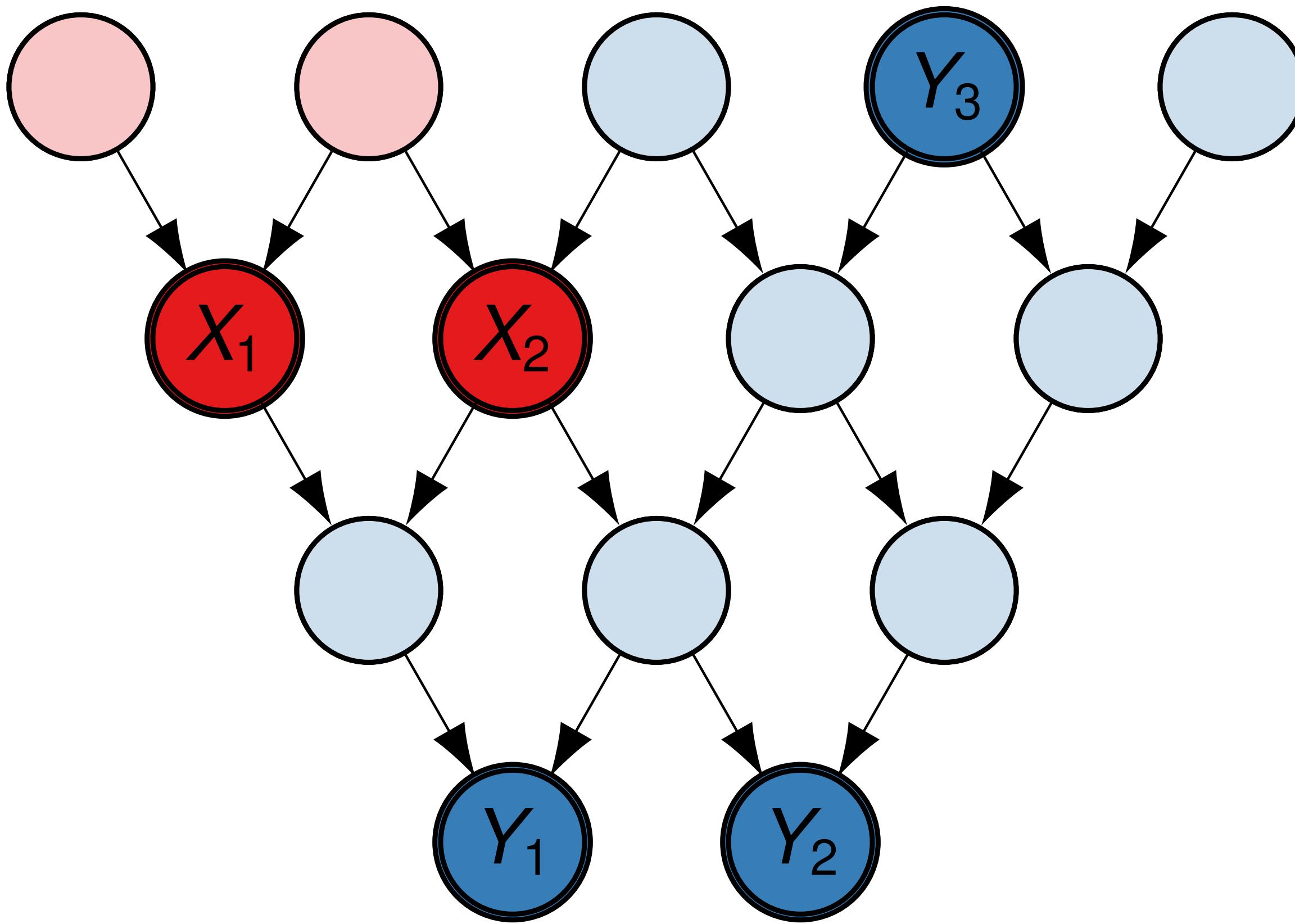
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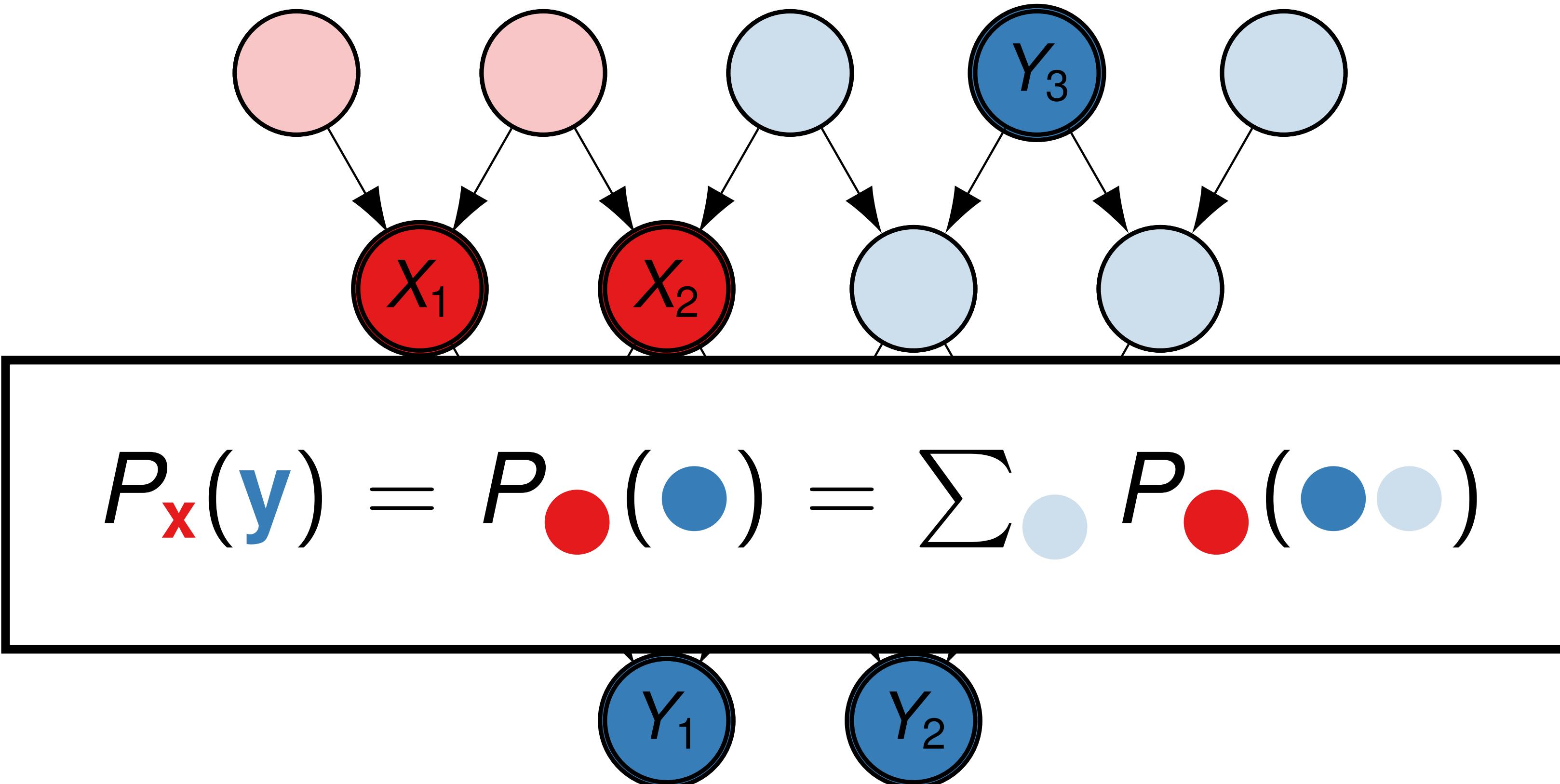
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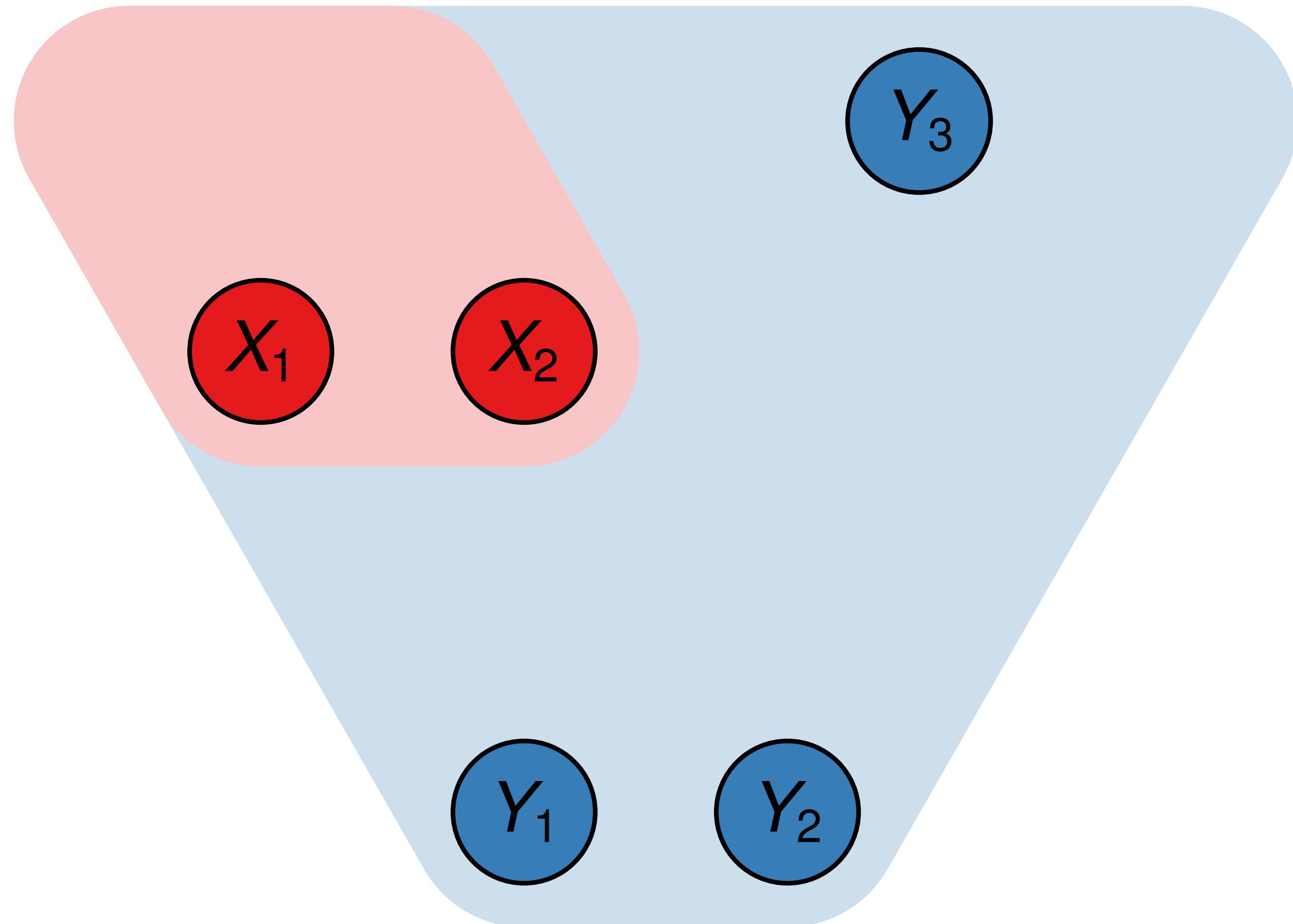
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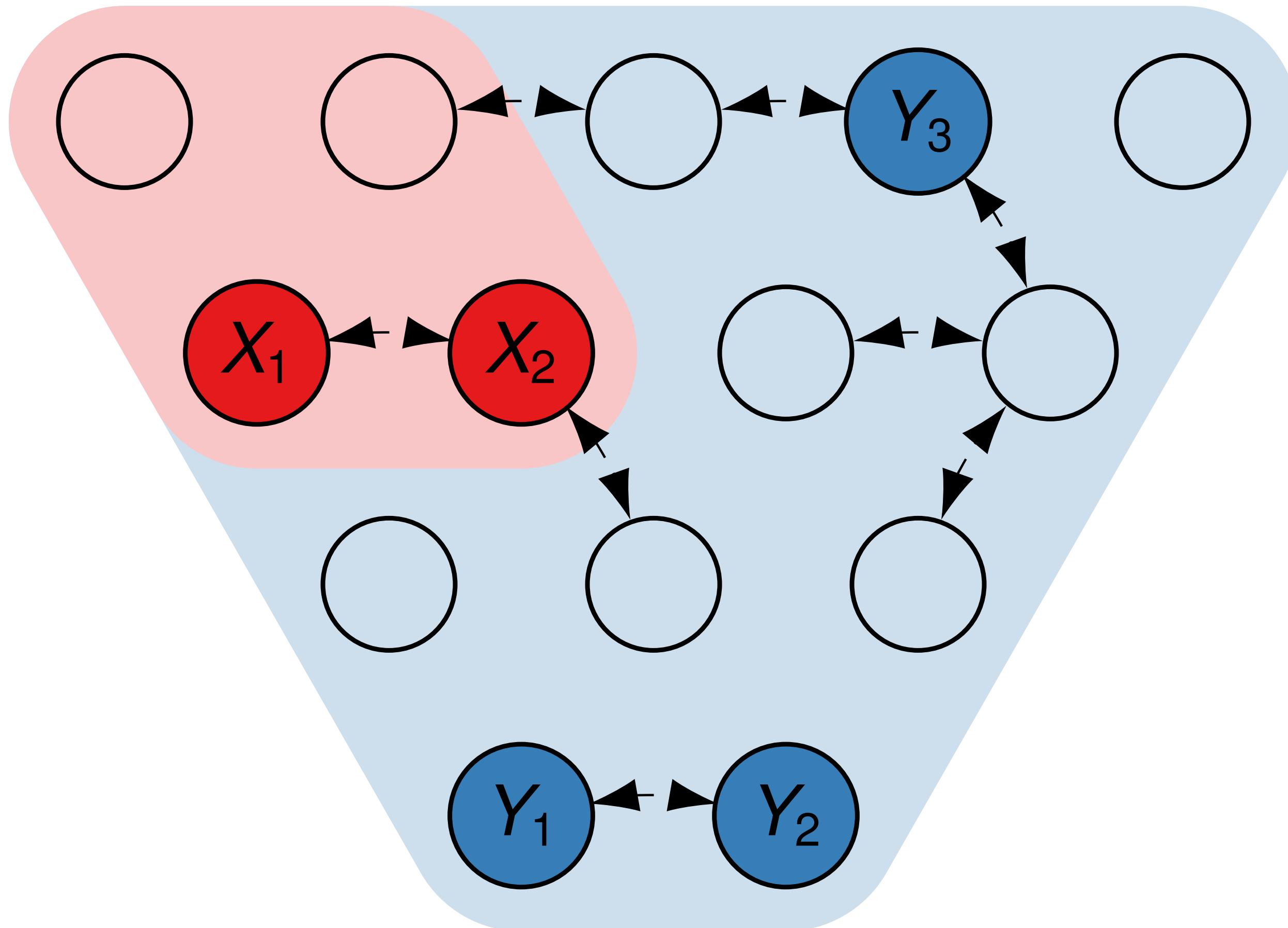
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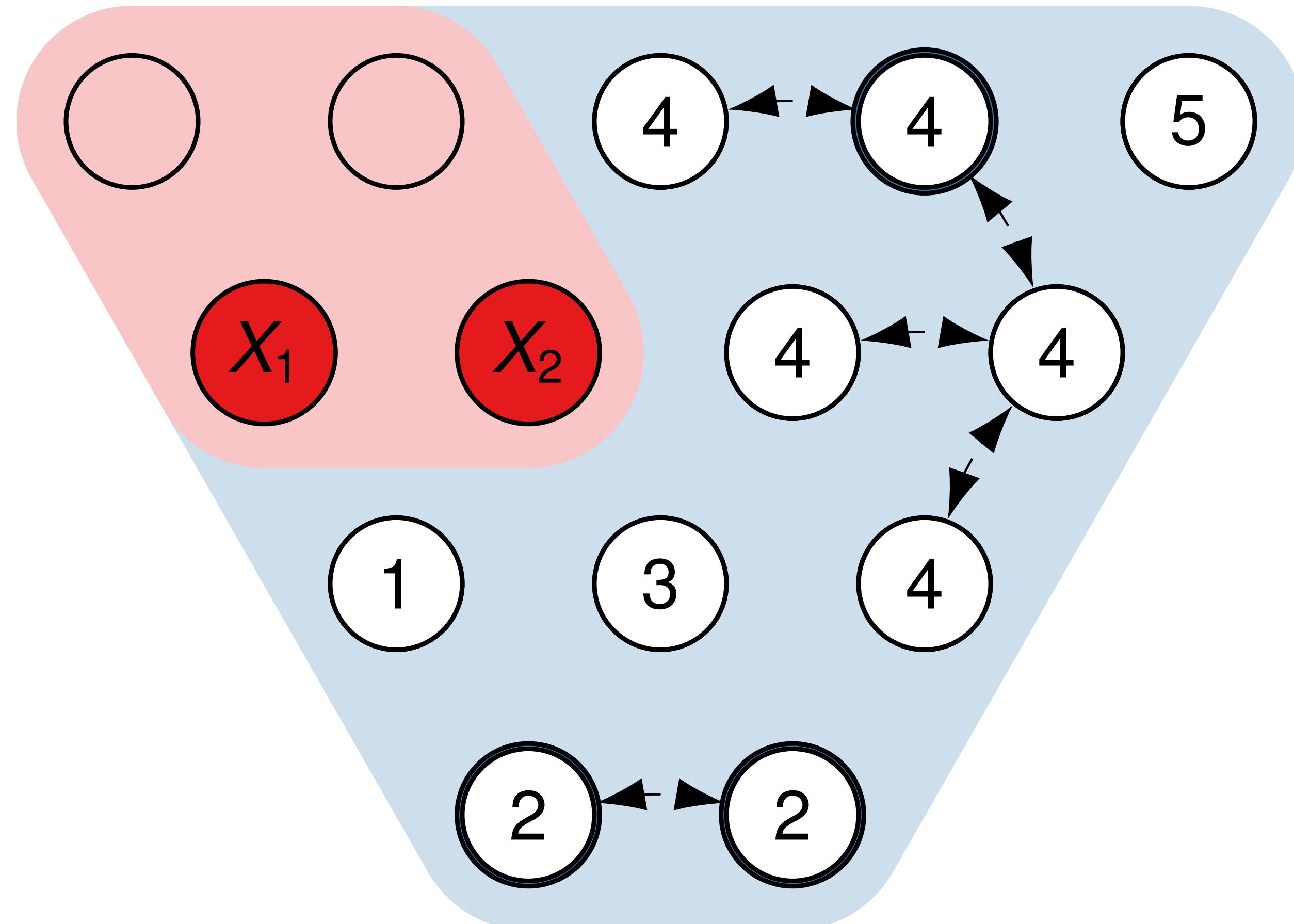
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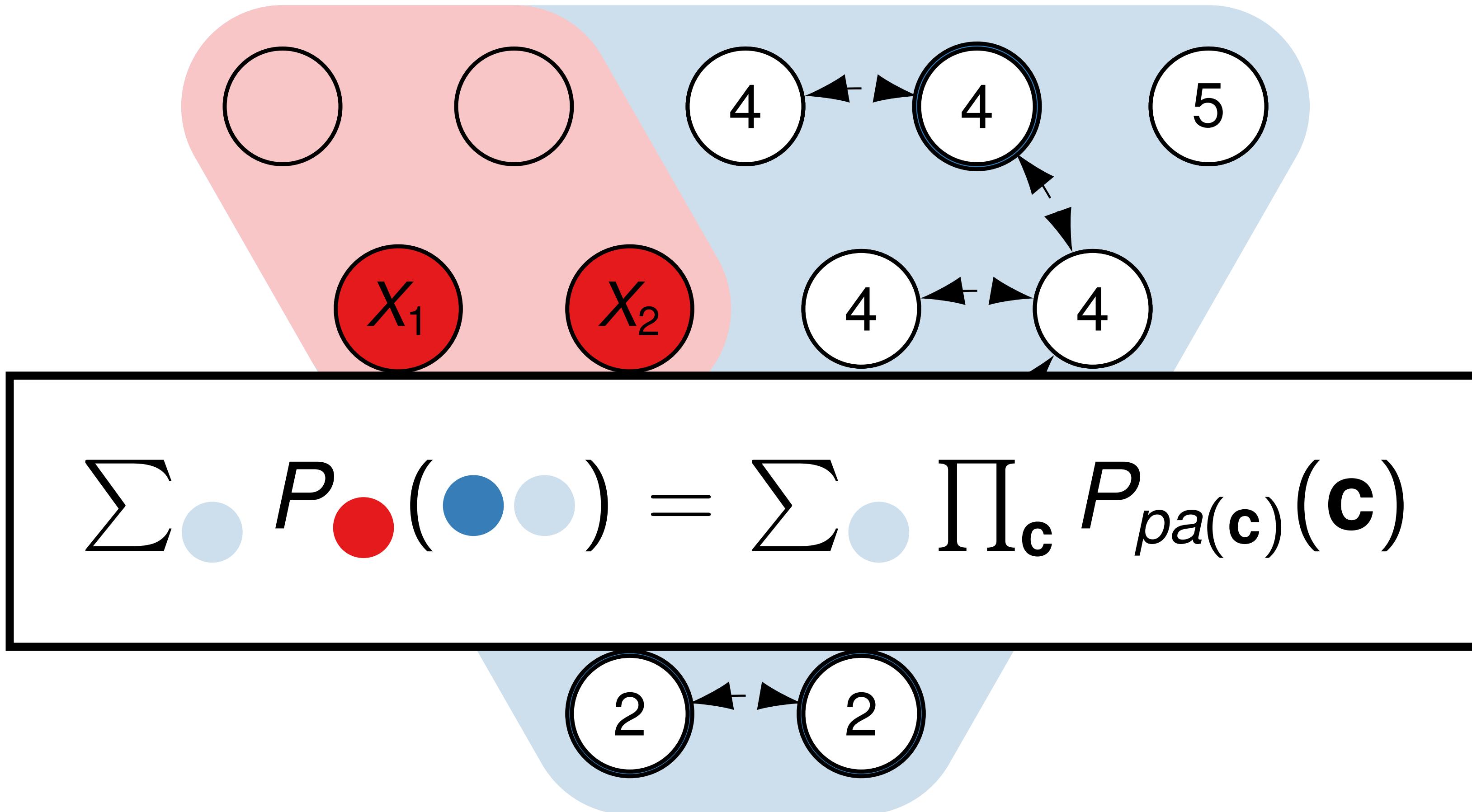
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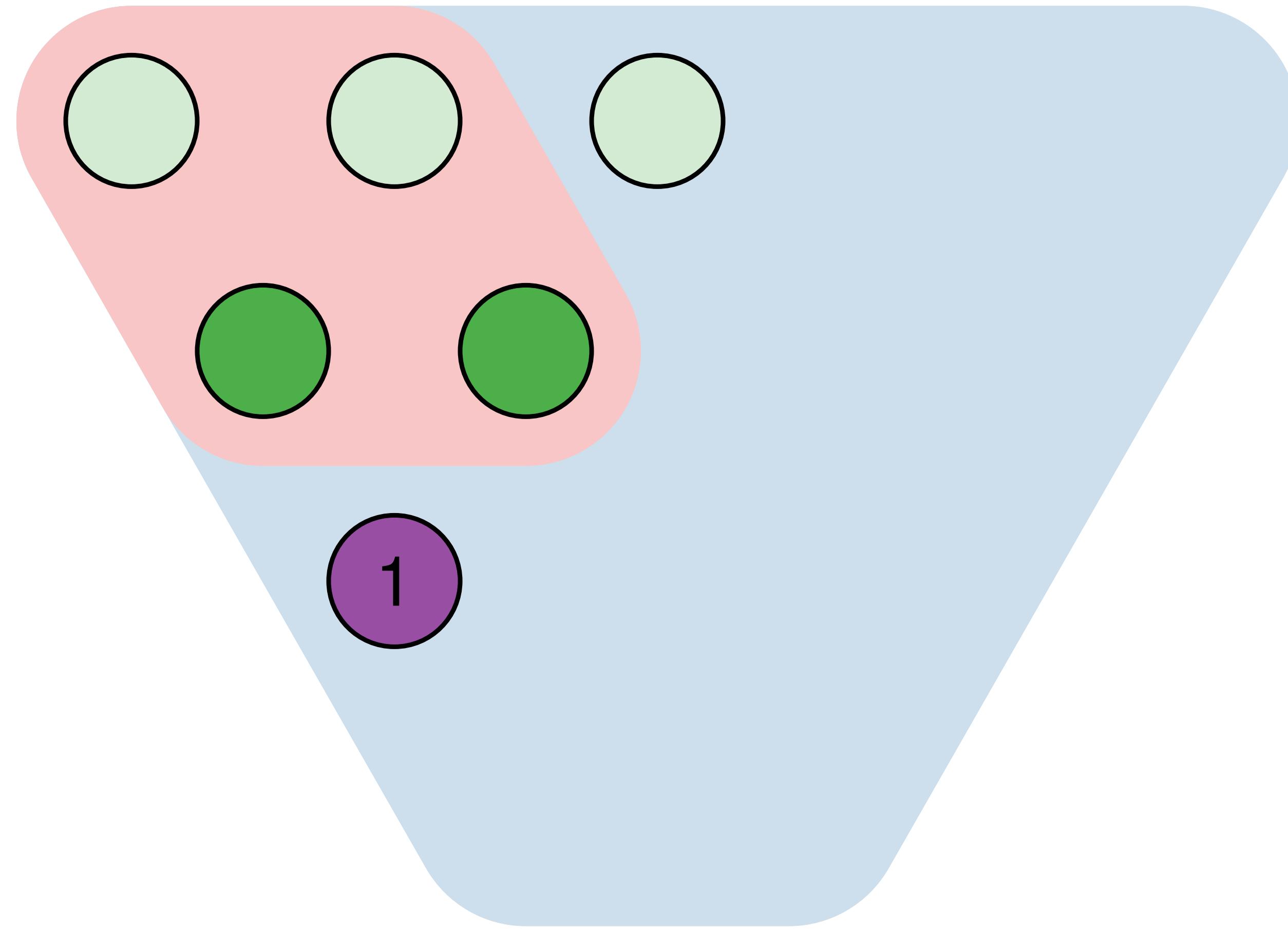
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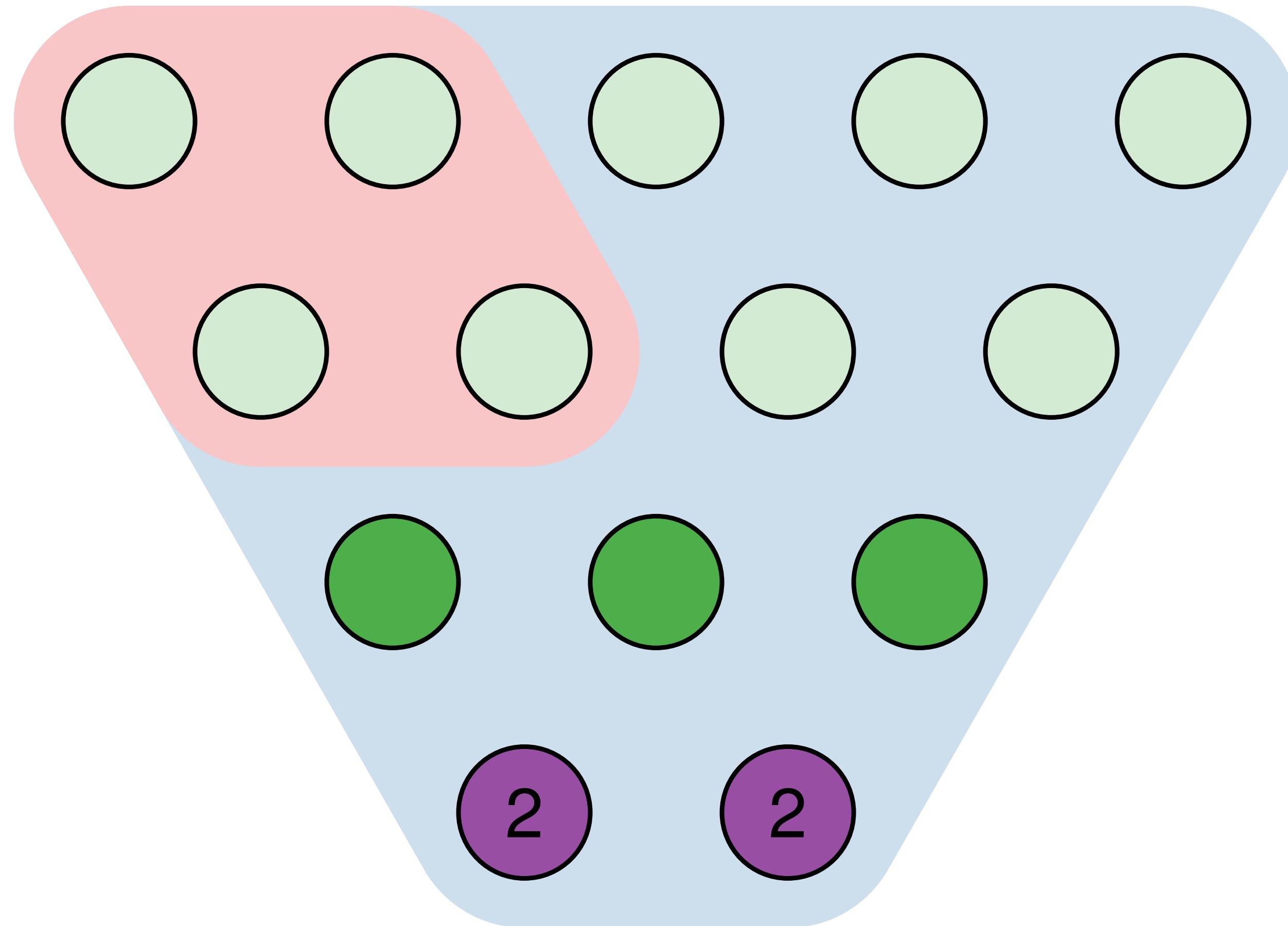
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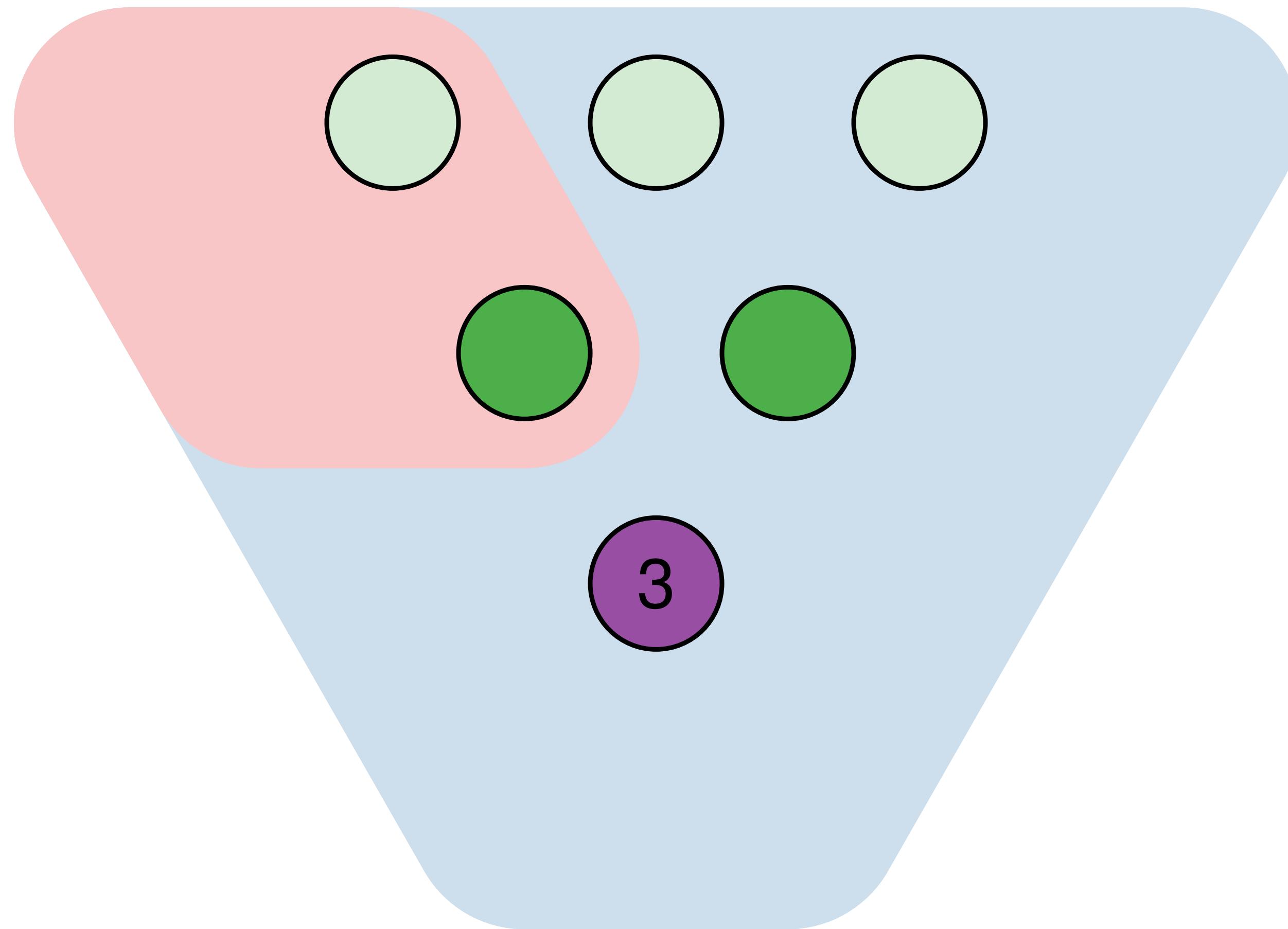
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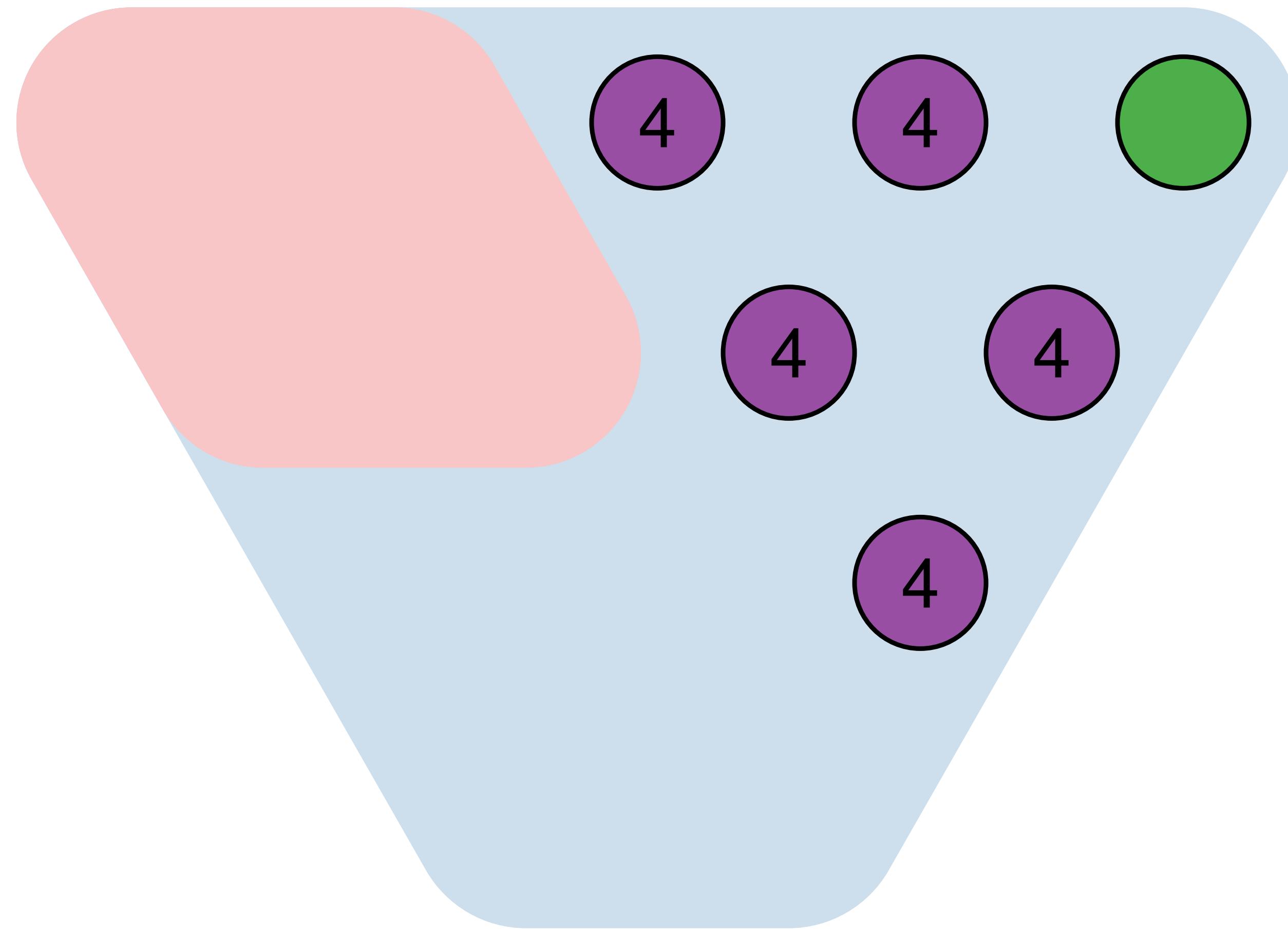
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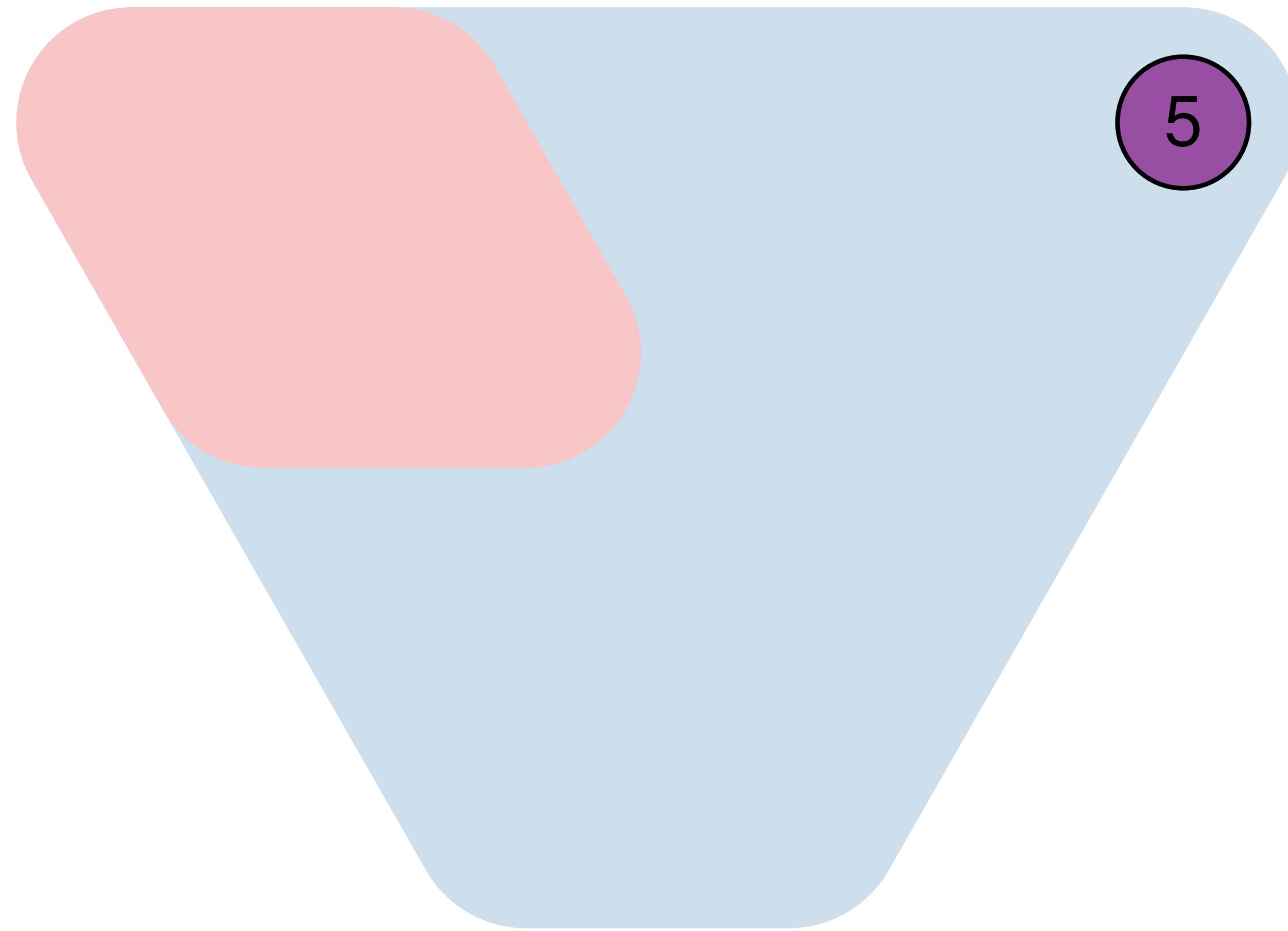
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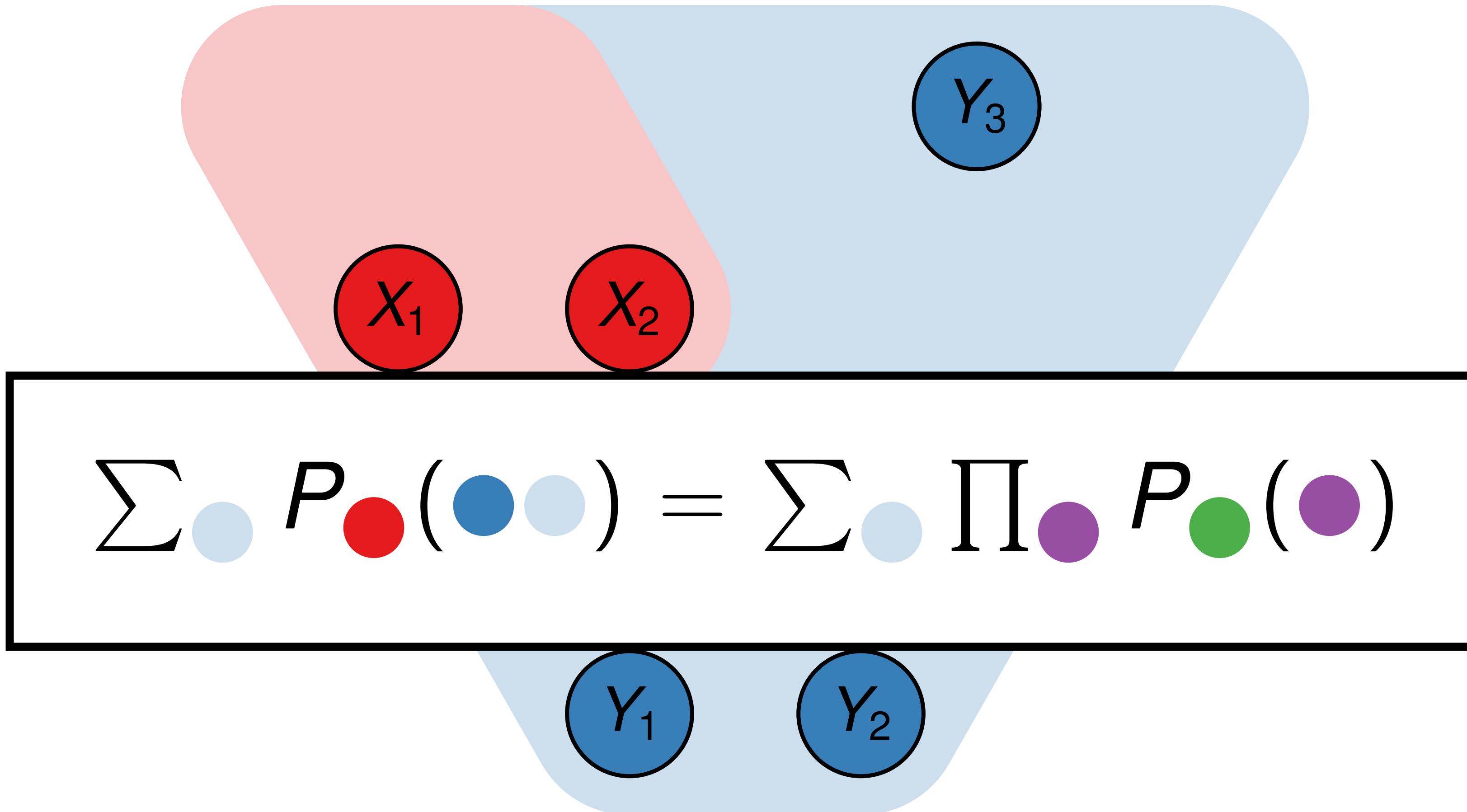
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# Algorithm for gID (1<sup>st</sup> phase)

```

function GID(, ,  $\mathcal{G}$ ,  $Z$ )
  if  $\exists_{z \in Z} z = Z \cap V$  then                                 $\triangleright$  check whether a matching experiment exists
    return  $P_{z \setminus V, \textcolor{red}{\bullet}}(\bullet)$ 

  if  $V \neq An(\bullet)_{\mathcal{G}}$  then                                 $\triangleright$  retain only the ancestors of 
    return GID(,   $\cap An(\bullet)_{\mathcal{G}}$ ,  $\mathcal{G}[An(\bullet)_{\mathcal{G}}], Z$ )

  if  $(W \leftarrow (V \setminus \bullet) \setminus An(\bullet)_{\mathcal{G}}) \neq \emptyset$  then           $\triangleright$  modify to a maximal intervention
    return GID(,   $\cup$  ,  $\mathcal{G}, Z$ )

   $S \leftarrow \mathcal{C}(\mathcal{G} \setminus \bullet \bullet)$ 
  if  $|S| > 1$  then                                 $\triangleright$  factorize into subqueries
    return  $\sum_{\bullet \in S} \prod_{\bullet \in S} GID(\bullet, \bullet \bullet, \mathcal{G}, Z)$ 

  for  $Z \in \mathbb{Z}$  such that  $Z \cap V \subseteq \bullet \bullet$  do       $\triangleright$  identify with each of available distribution
    return SUB-ID(,   $\setminus Z, P_{(z \setminus V), \bullet \bullet \cap Z}, \mathcal{G} \setminus (Z \cap \bullet \bullet)$ ) if not NONE

  throw FAIL

```

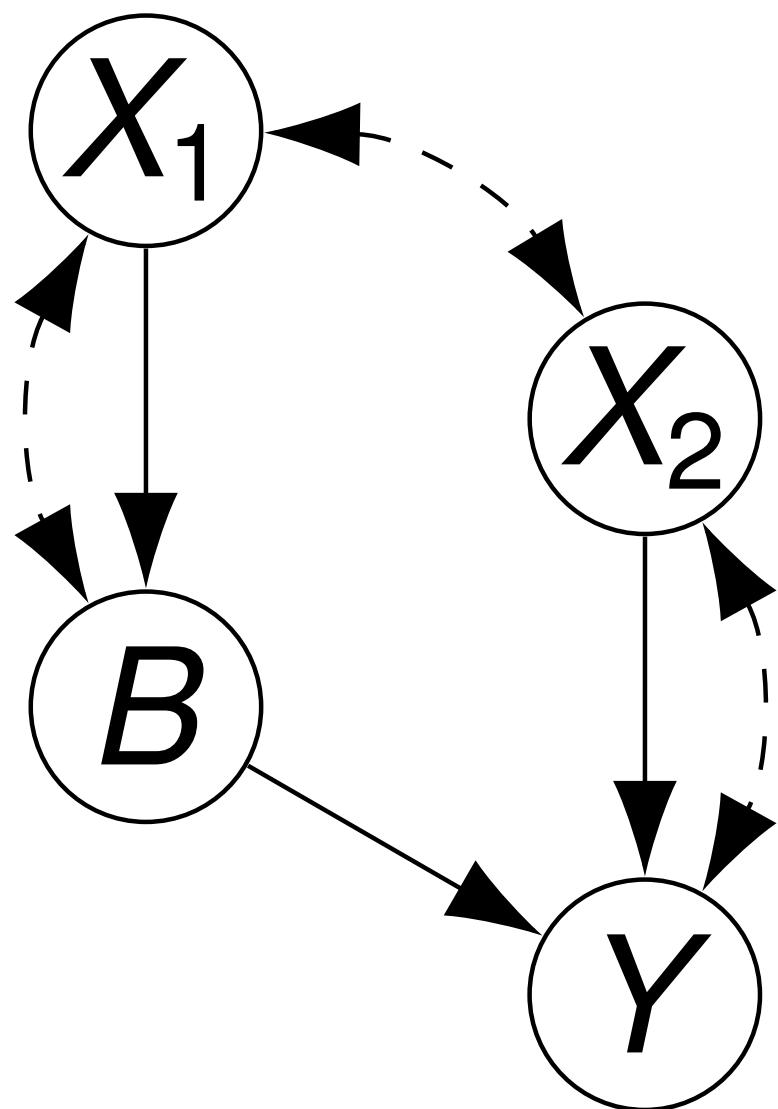
**sub-ID** is a simplified **ID** algorithm [SP'06], which returns **NONE** if failed.

# Algorithm for gID (2<sup>nd</sup> phase)

```
function SUB-ID(, , Q, G)
  {S} ← C(G \ 
  if  = ∅ then                                ▷ check identified
    return  $\sum_{v \setminus \bullet} Q(v)$ 
  if V ≠ An()G then                                    ▷ retain only the ancestors of 
    return SUB-ID(,  ∩ An()G,  $\sum_{v \setminus An(\bullet)_G} Q, G[An(\bullet)_G]$ )
  if C(G) = V then                            ▷ check the existence of a hedge
    return NONE
  if S ∈ C(G) then                           ▷ check identifiable
    return  $\sum_{s \setminus \bullet} \prod_{v_i \in \bullet} Q(v_i | v_\pi^{(i-1)})$ .
  if S ⊂neq; S' ∈ C(G) then                ▷ modify input (query, distribution, and graph)
    return SUB-ID(
```

# Example: Drug-Drug Interactions

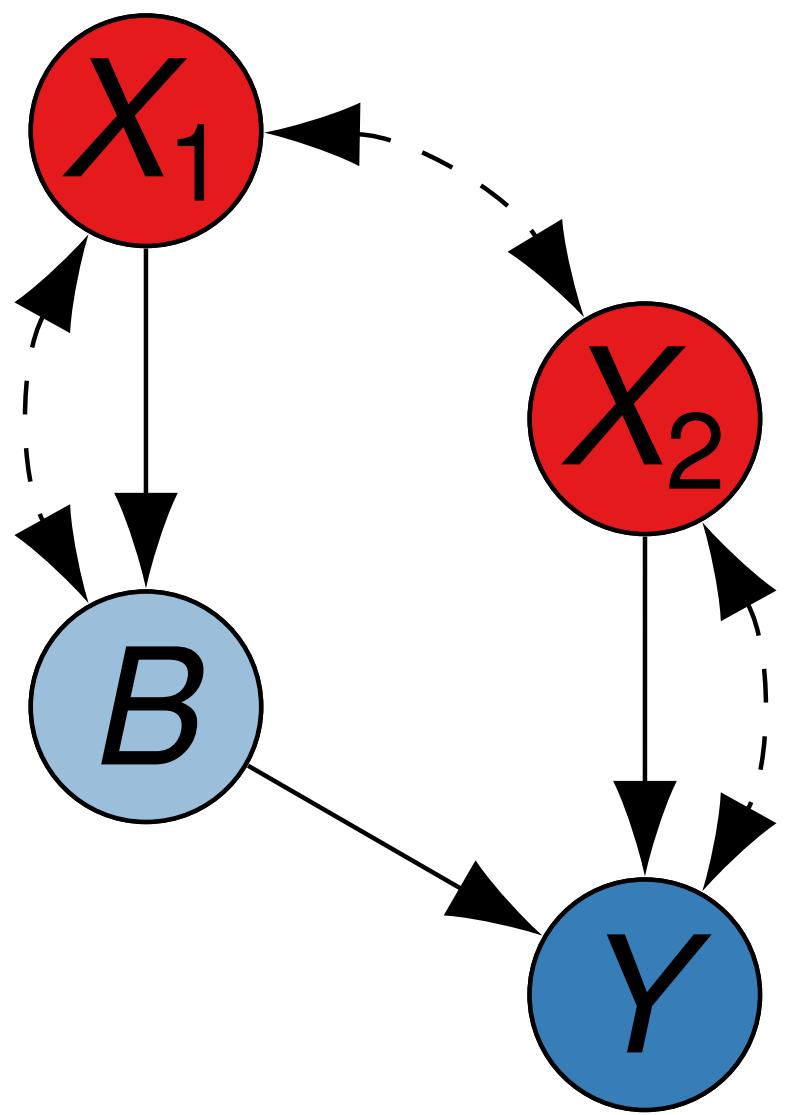
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$$P_{x_1, x_2}(y)$$

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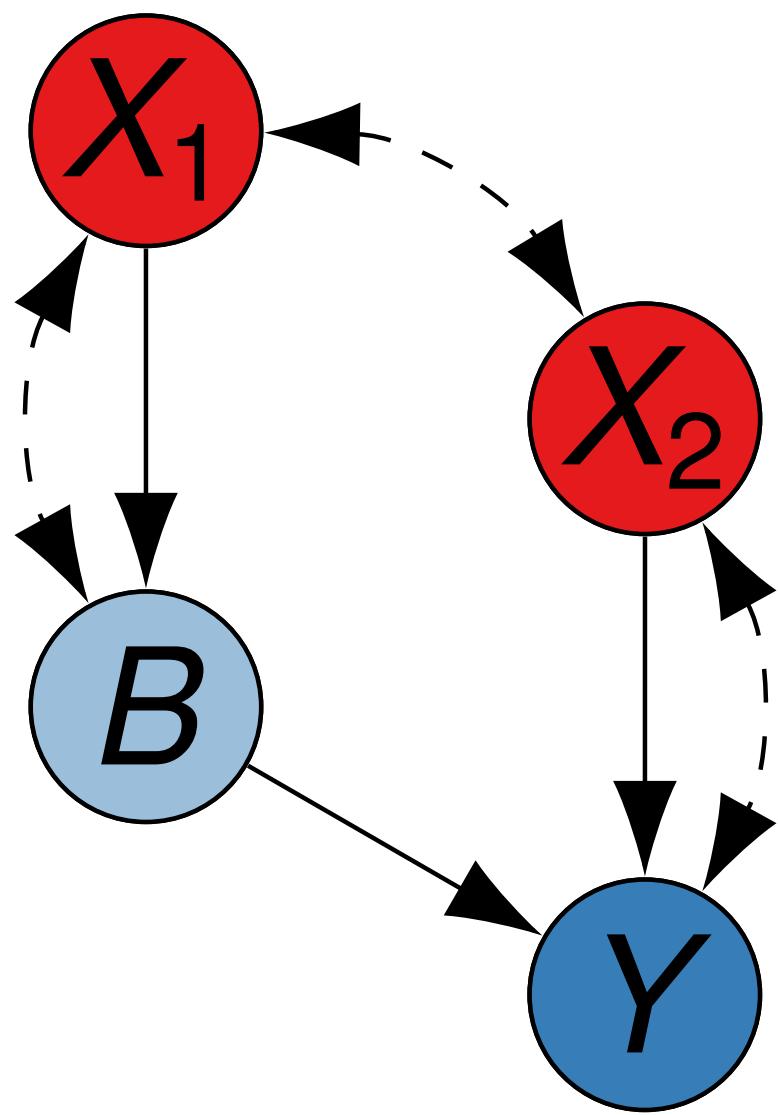
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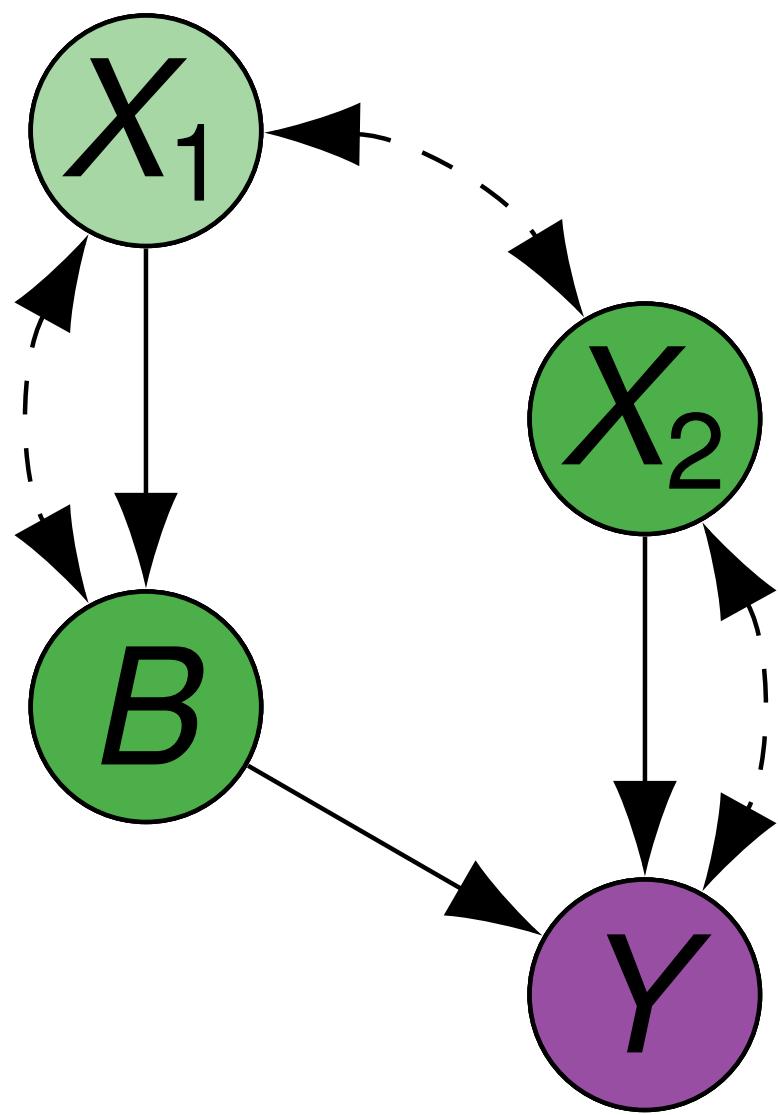
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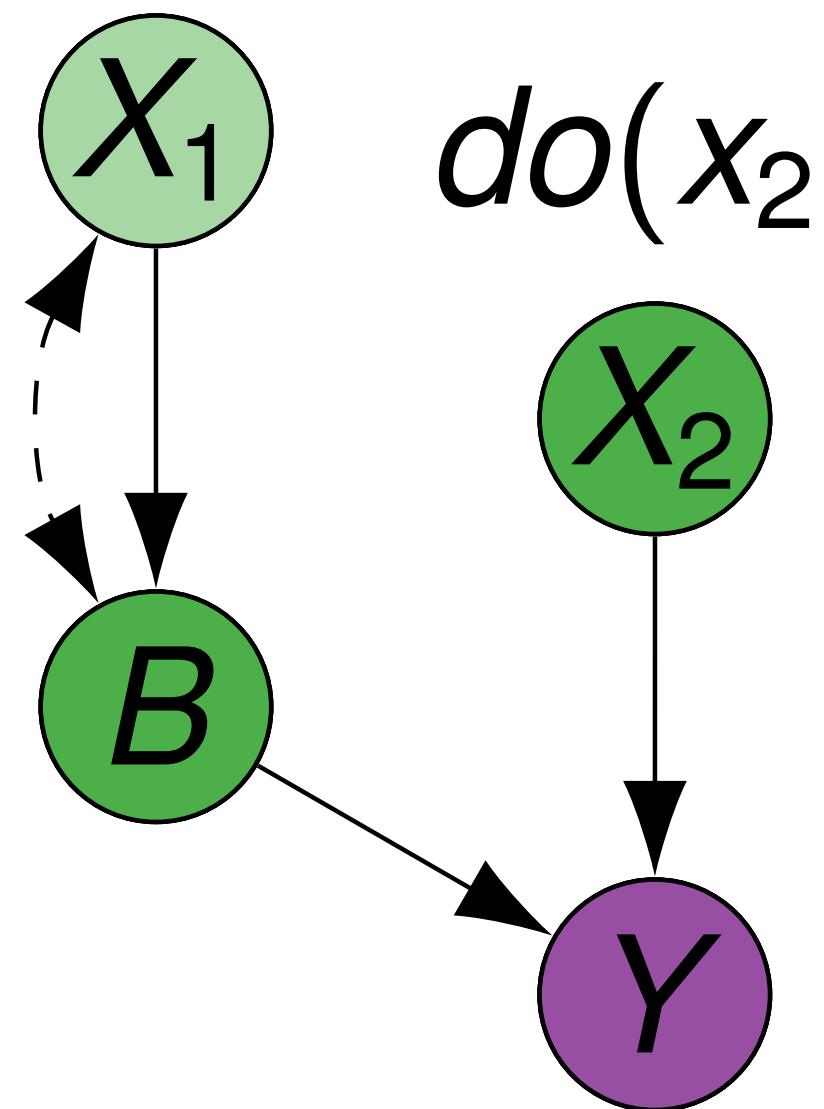
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## non-g-identifiability

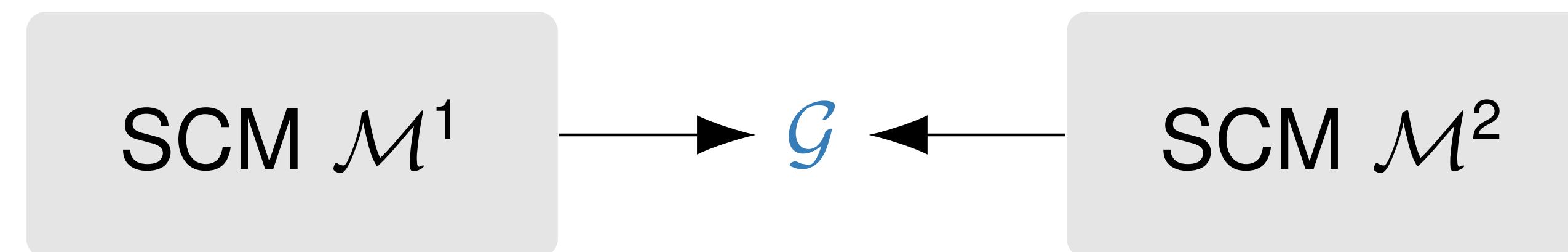
– the failure condition & a prohibiting structure

# Proving Non-g-Identifiability

To prove  $P_x(y)$  is **not g-identifiable** from  $\mathbb{Z}$  in  $\mathcal{G}$ ,

# Proving Non-g-Identifiability

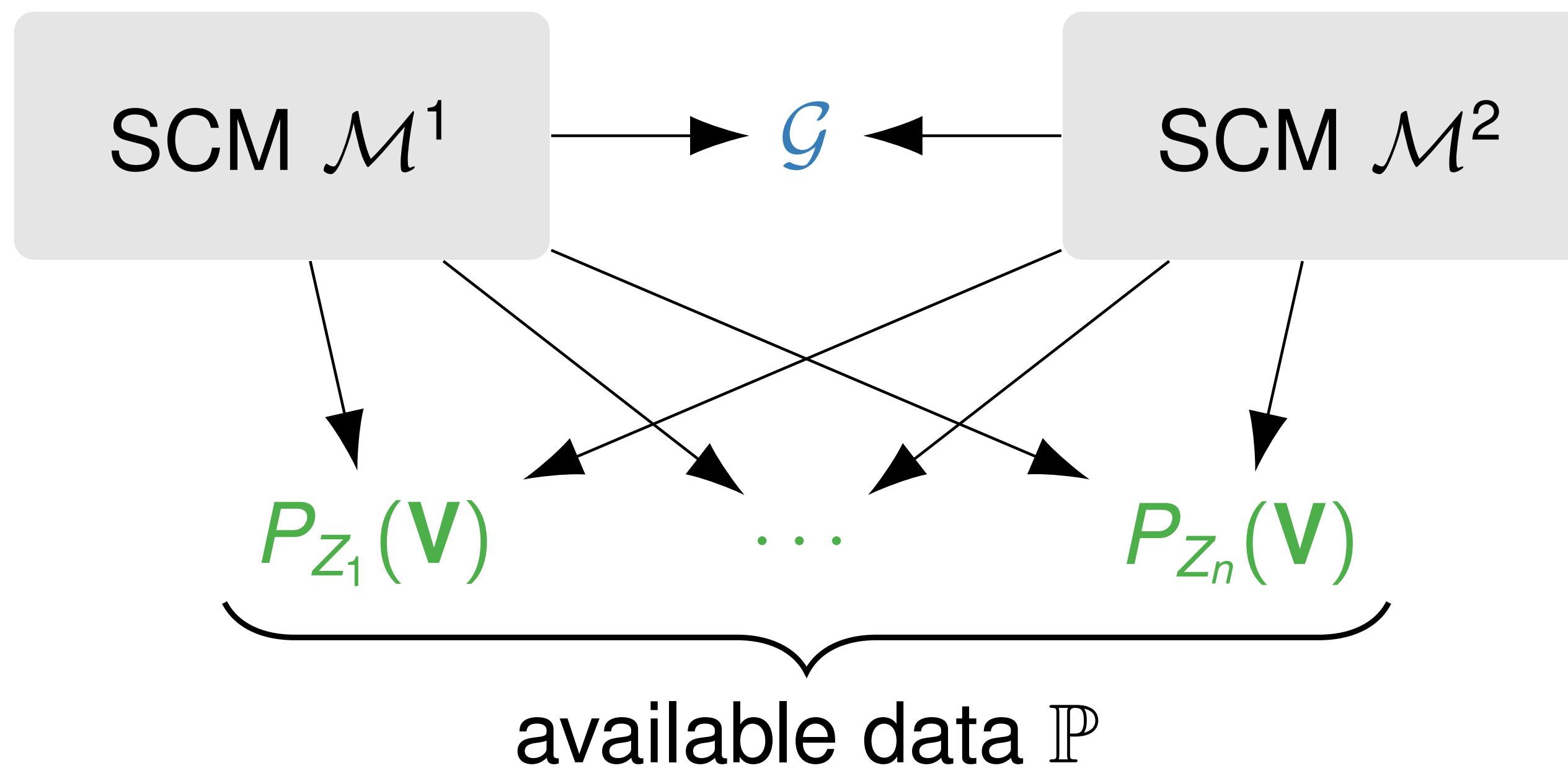
To prove  $P_x(y)$  is **not g-identifiable** from  $\mathbb{Z}$  in  $\mathcal{G}$ , we construct two causal models  $\mathcal{M}_1$  and  $\mathcal{M}_2$  compatible with  $\mathcal{G}$



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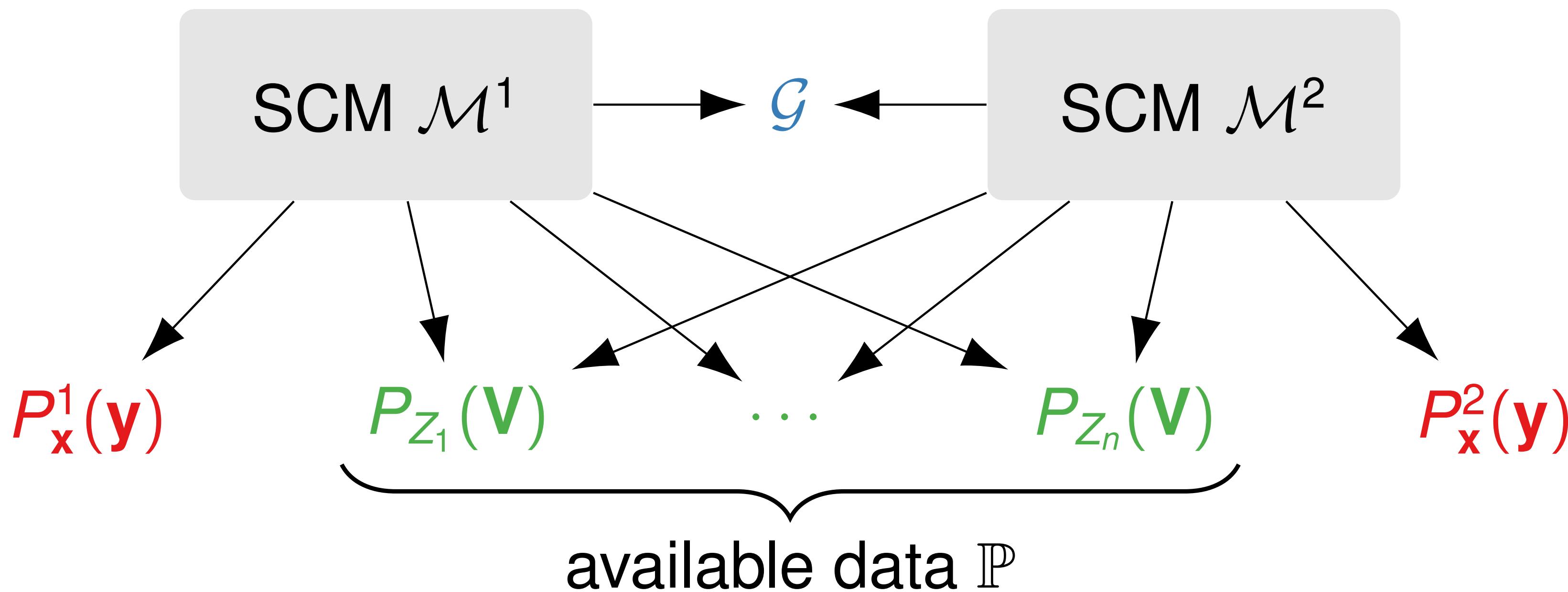
$$P_{\mathbf{z}}^1(\mathbf{v}) = P_{\mathbf{z}}^2(\mathbf{v}) \text{ for all } \mathbf{Z} \in \mathbb{Z}, \mathbf{z} \in \mathfrak{X}_{\mathbf{z}},$$



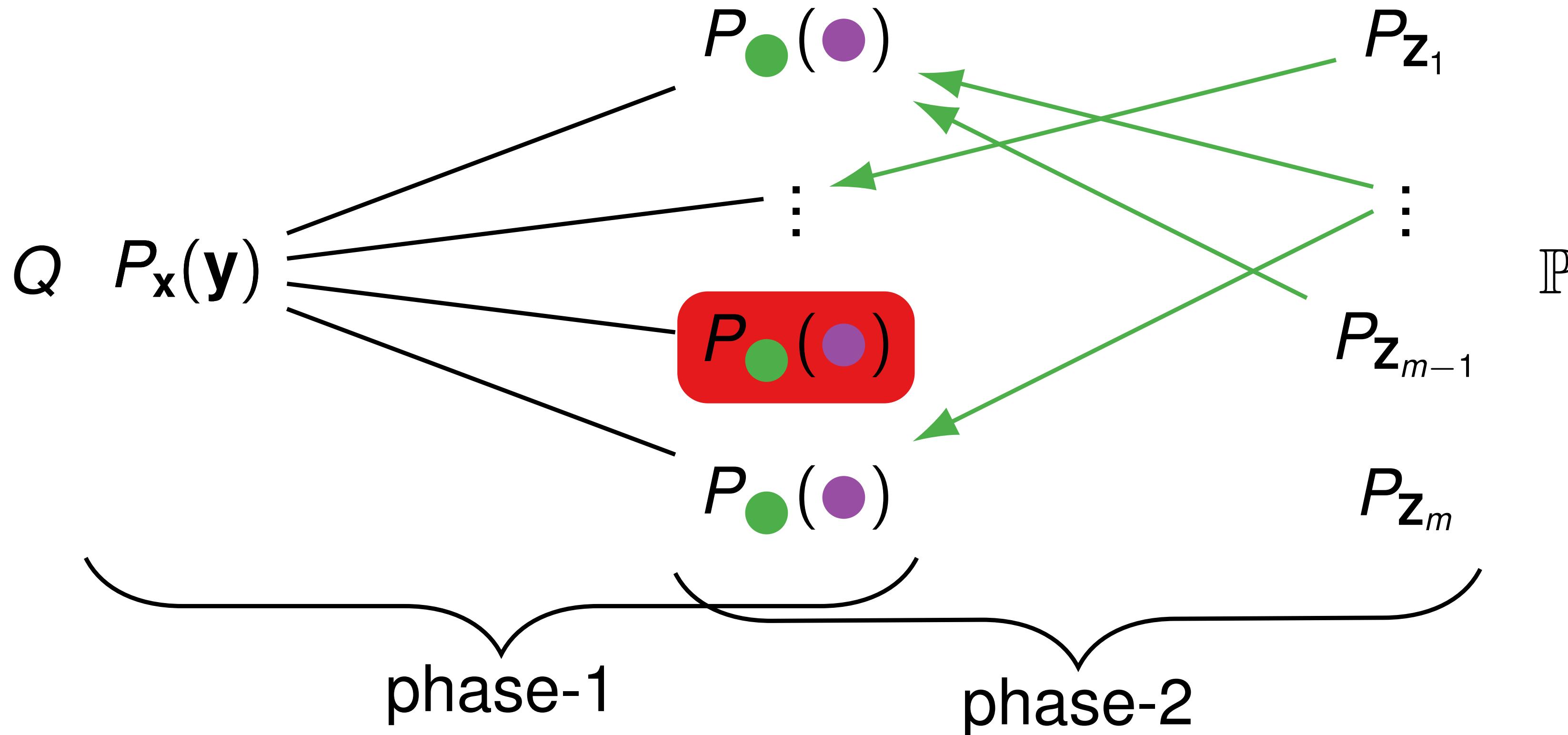
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$$P_z^1(v) = P_z^2(v) \text{ for all } z \in \mathbb{Z}, z \in \mathfrak{X}_z, \text{ but } P_x^1(y) \neq P_x^2(y).$$



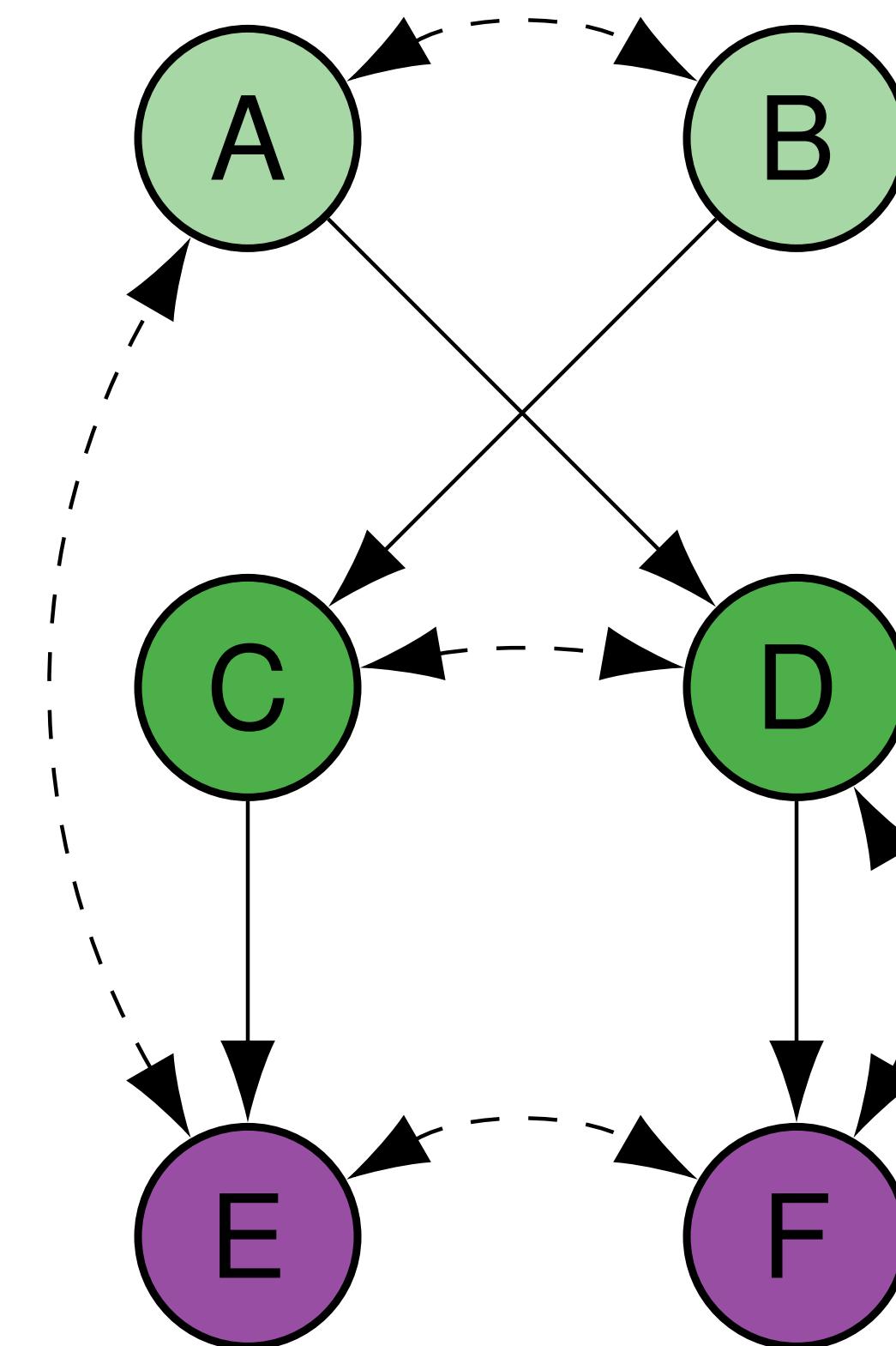
# Recall the failed factor ...



$\exists P_z_i(\bullet)$  such that  $\forall P_z_j \in \mathbb{P}$  fails

# $P_{\bullet}(\bullet)$ versus $P_{z_i} \in \mathbb{P}$ (phase-2)

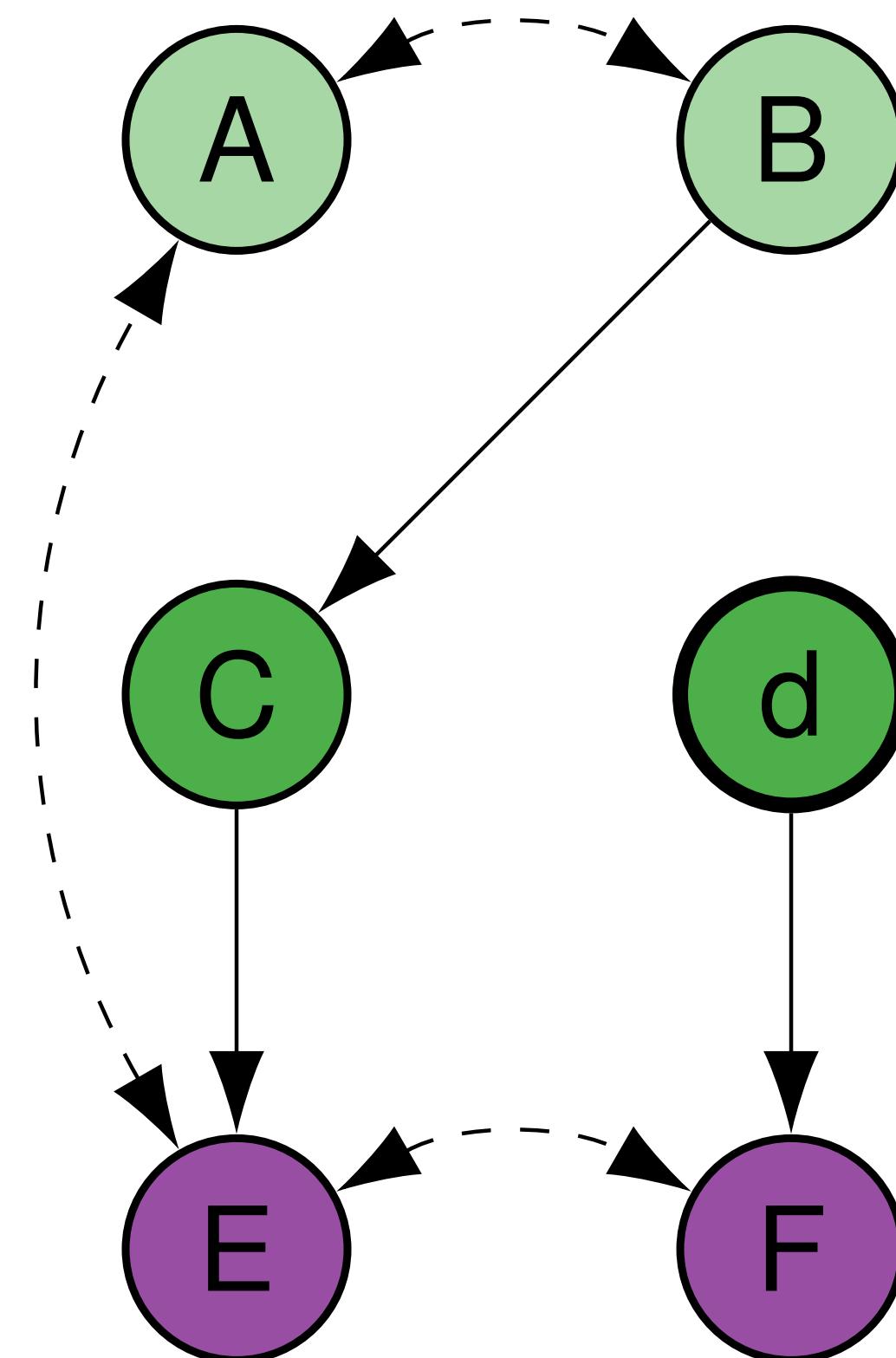
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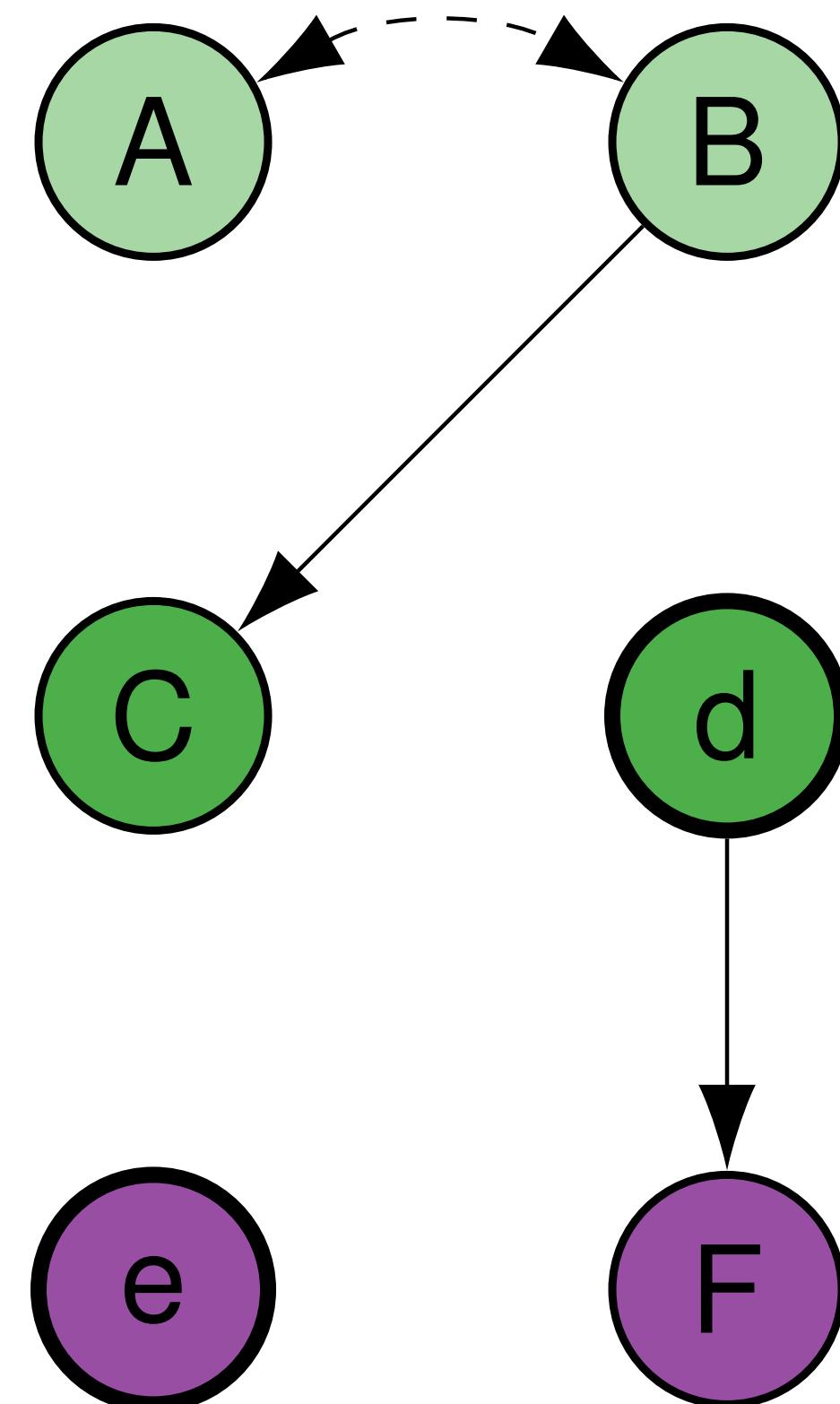


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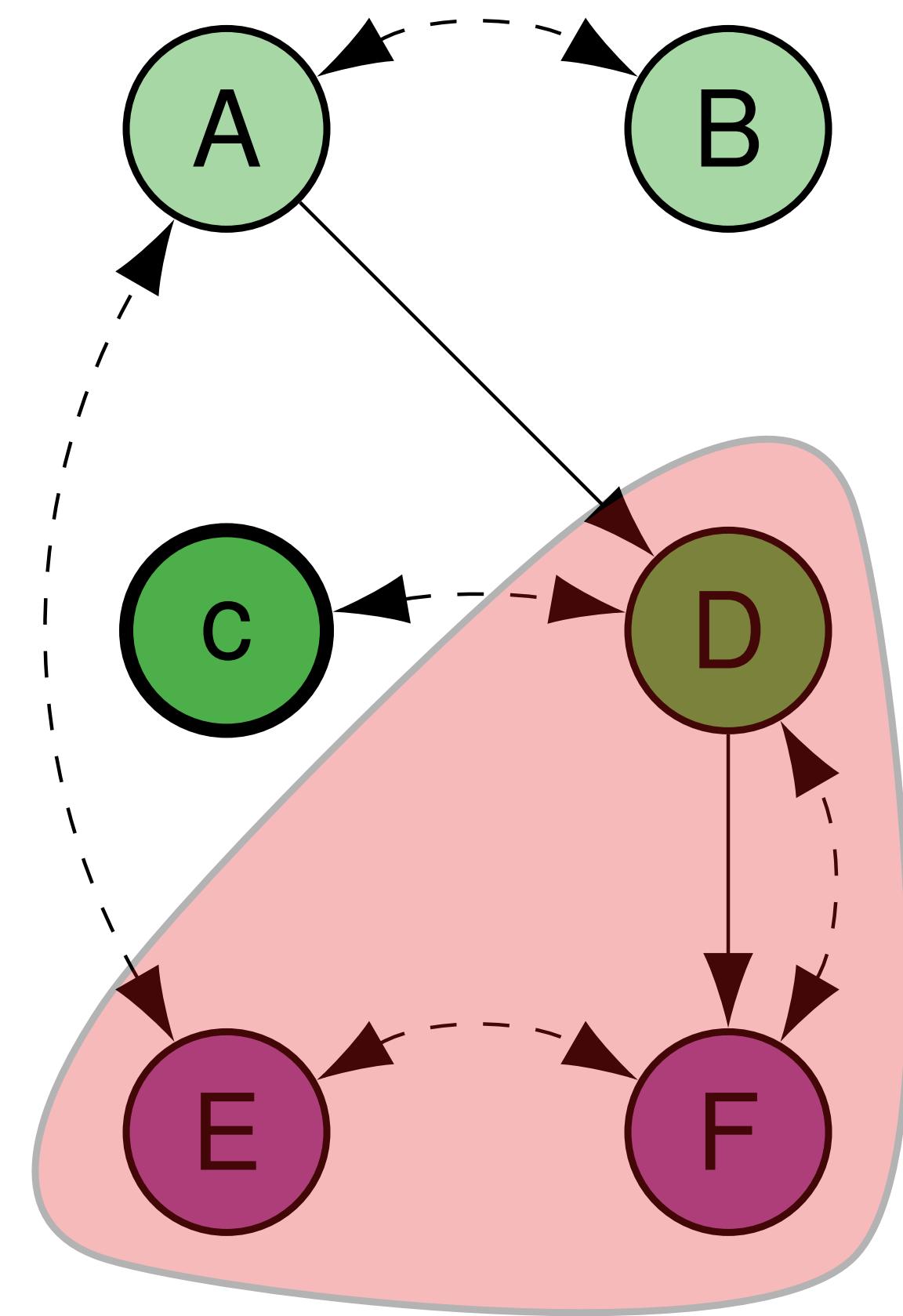


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- ✗ (the **bad**)  $\exists$  hedge [SP'06],  
e.g.,  $P_C$



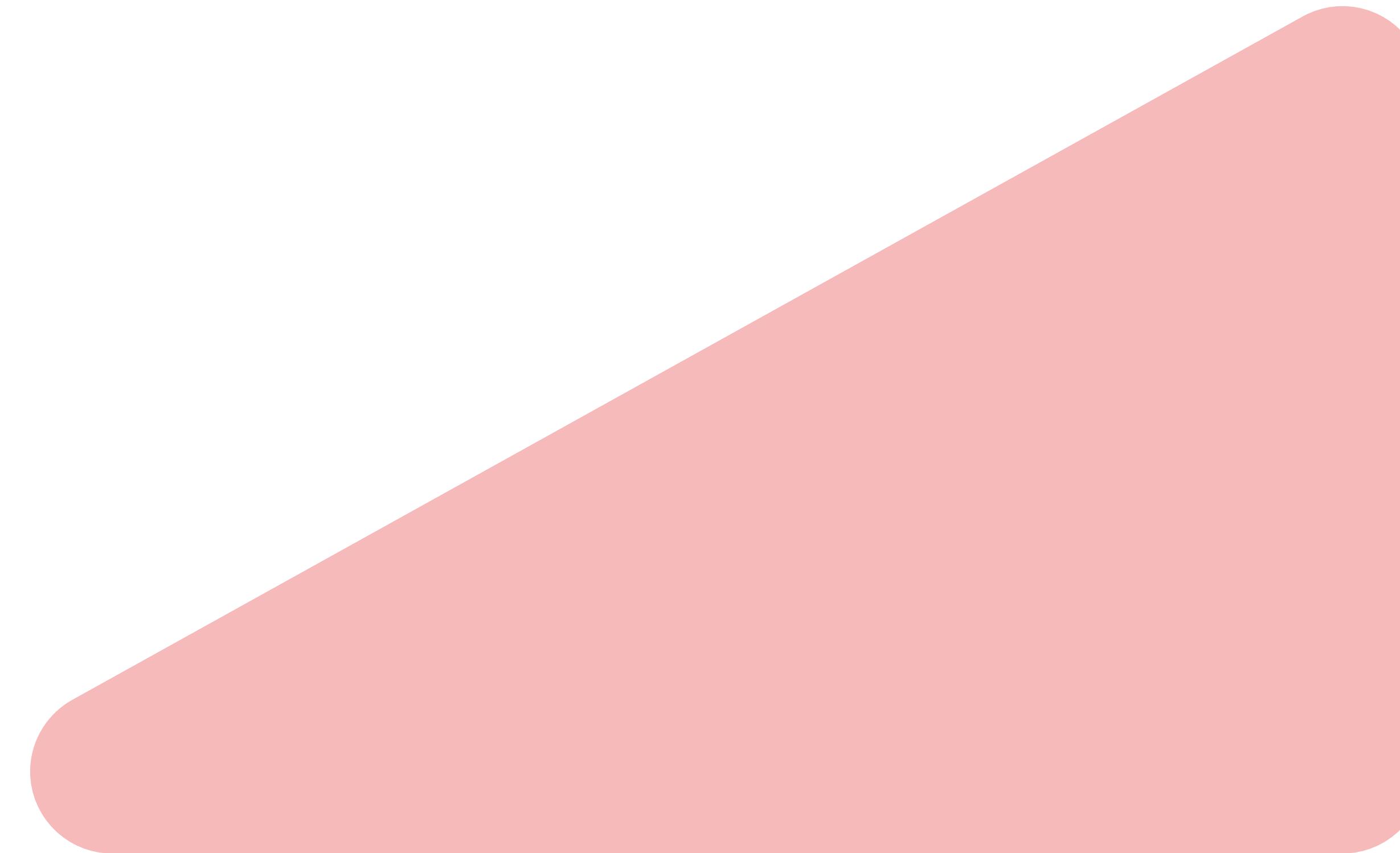
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# Failure $\Rightarrow \exists$ Thicket

- A **thicket** is the superimposition of **hedges** (the bad structure).
- (if every experiment intersects with ●, confounded ●s form a ‘degenerate **thicket**.’)

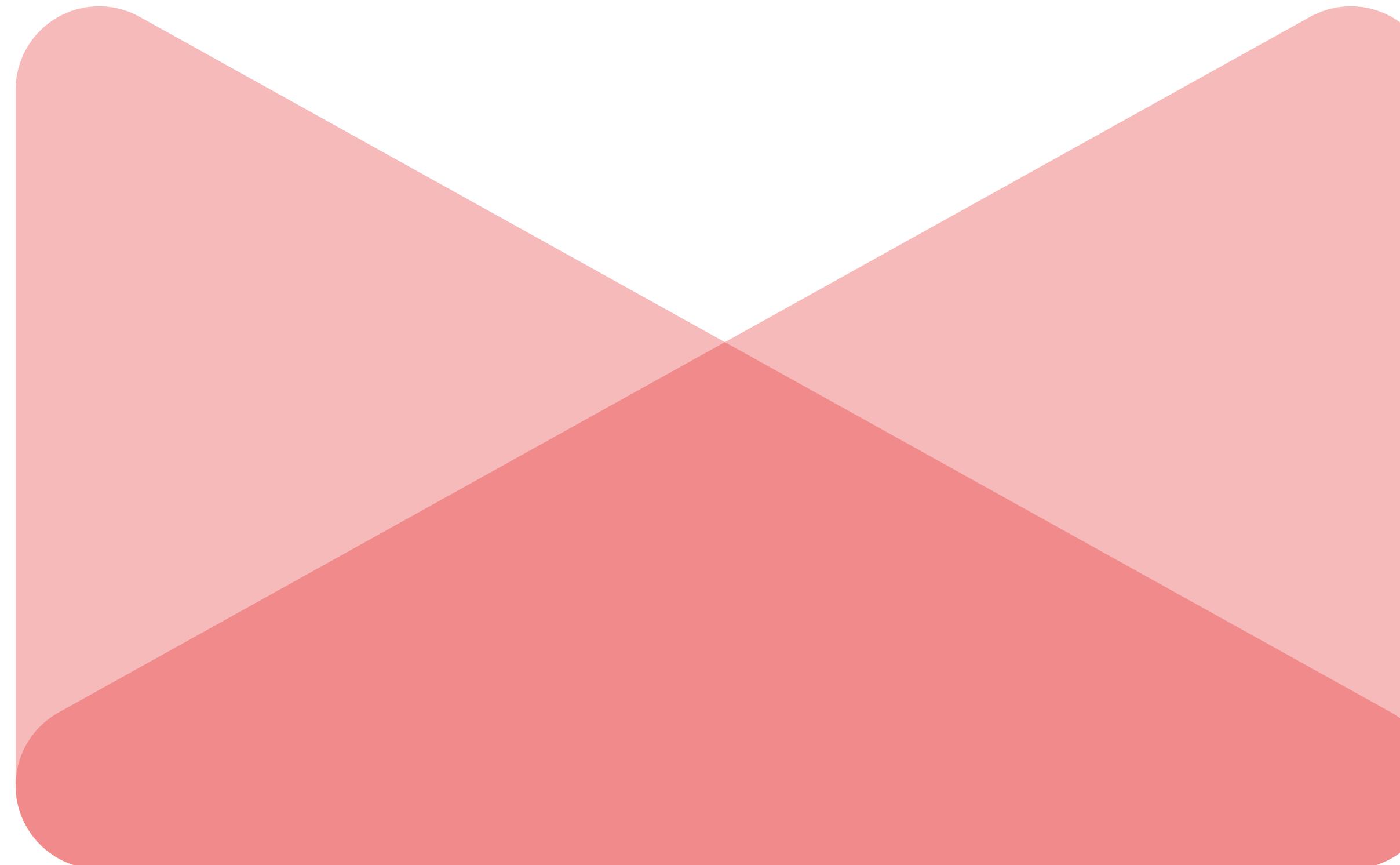
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## Definition

Let  $\mathbf{R}$  be a non-empty set of variables and  $\mathbb{Z}$  be a collection of sets of variables in  $\mathcal{G}$ . A thicket  $\mathcal{J} \subseteq \mathcal{G}$  is an  $\mathbf{R}$ -rooted c-component consisting of a minimal c-component over  $\mathbf{R}$  and hedges

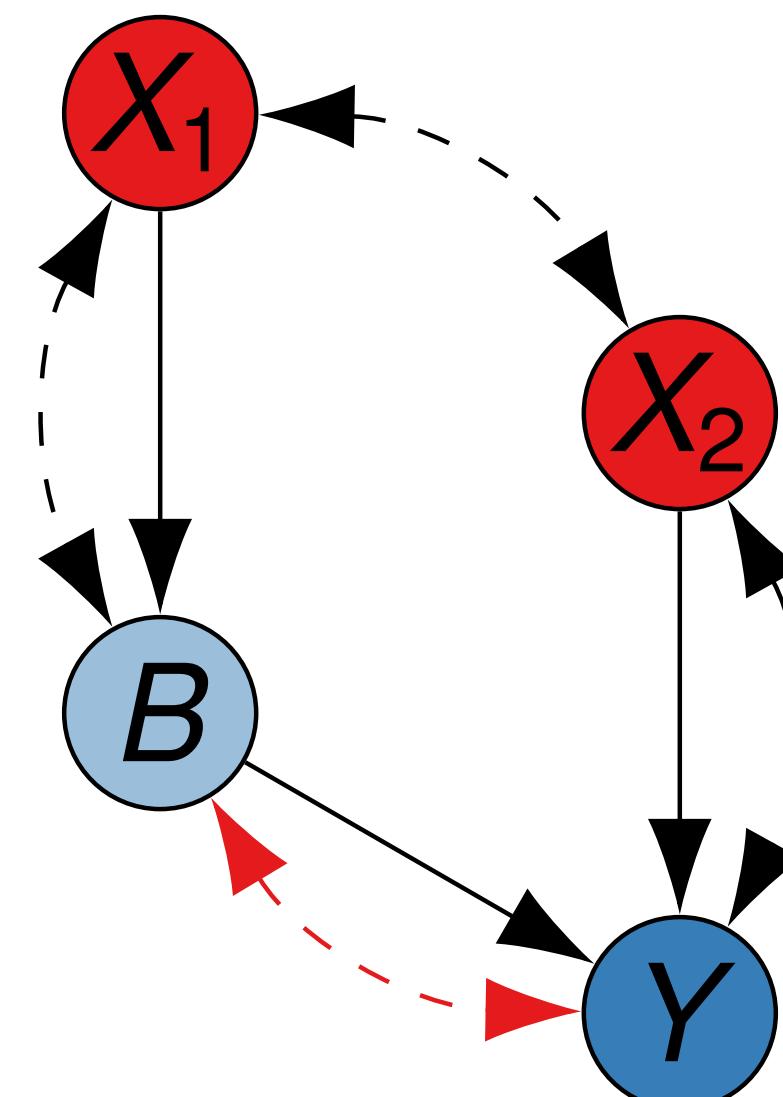
$$\mathbb{F}_{\mathcal{J}} = \{\langle \mathcal{F}_z, \mathcal{J}[\mathbf{R}] \rangle \mid \mathcal{F}_z \subseteq \mathcal{G} \setminus \mathbb{Z}, \mathbb{Z} \cap \mathbf{R} = \emptyset\}_{z \in \mathbb{Z}}.$$

Let  $\mathbf{X}, \mathbf{Y}$  be disjoint sets of variables in  $\mathcal{G}$ . A thicket  $\mathcal{J}$  is said to be formed for  $P_x(\mathbf{y})$  in  $\mathcal{G}$  with respect to  $\mathbb{Z}$  if  $\mathbf{R} \subseteq An(\mathbf{Y})_{\mathcal{G}_{\mathbf{X}}}$  and every hedgelet of each hedge  $\langle \mathcal{F}_z, \mathcal{J}[\mathbf{R}] \rangle$  intersects with  $\mathbf{X}$ .

# Thicket: Drug-Drug Interactions

Given  $Z = \{\{X_1\}, \{X_2\}\}$  with  $Q = P(y|do(x_1, x_2))$  (\*  $P_{\bullet}(\bullet) = P_{\bullet}(\bullet)$ )

- The query is **not g-identifiable**, see  $B \leftarrow \_ \rightarrow Y$

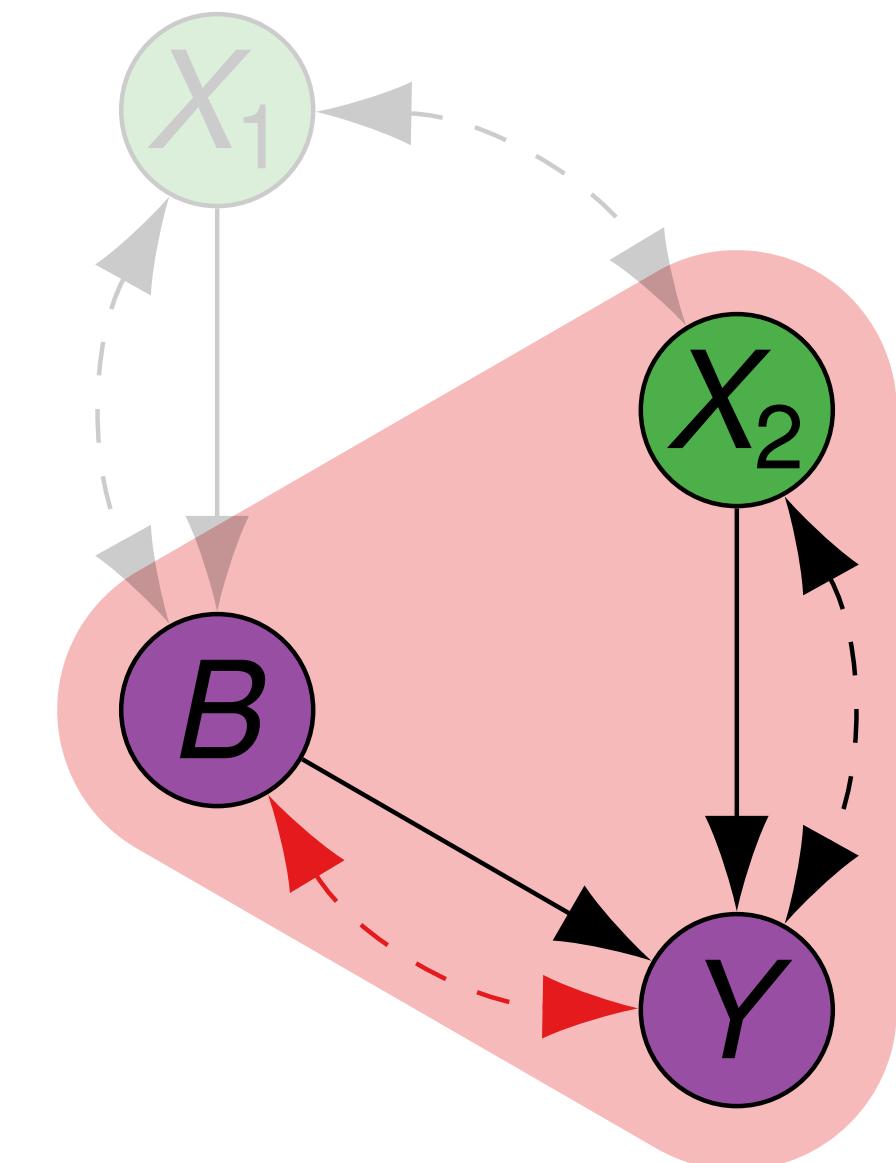


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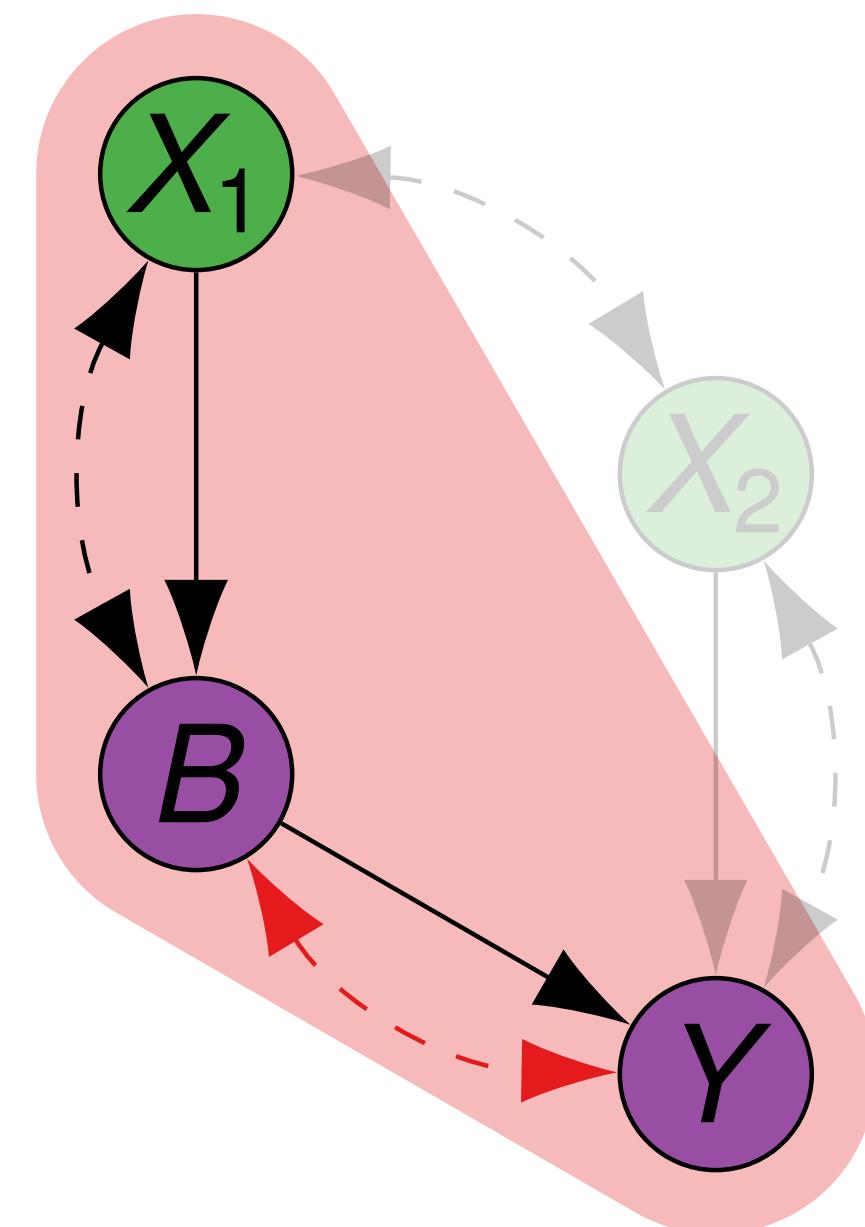


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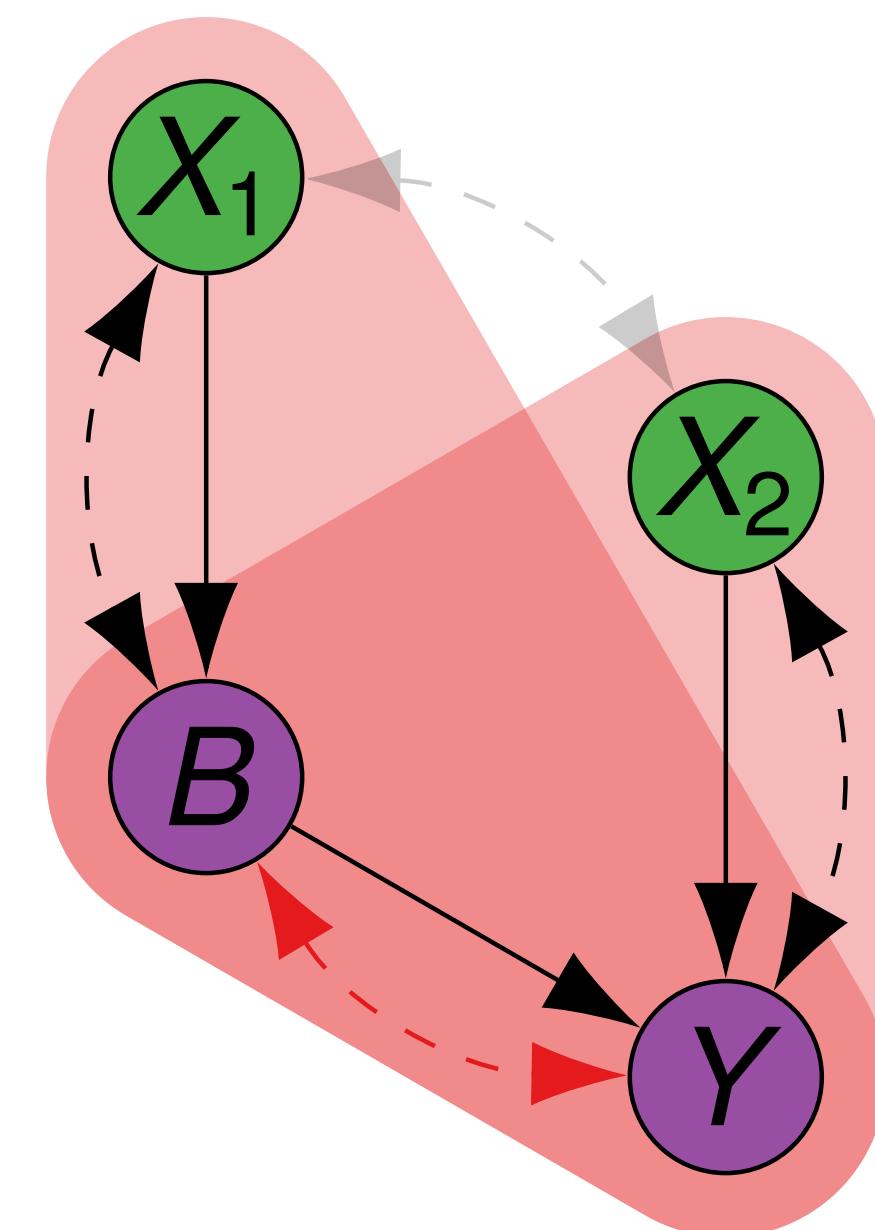


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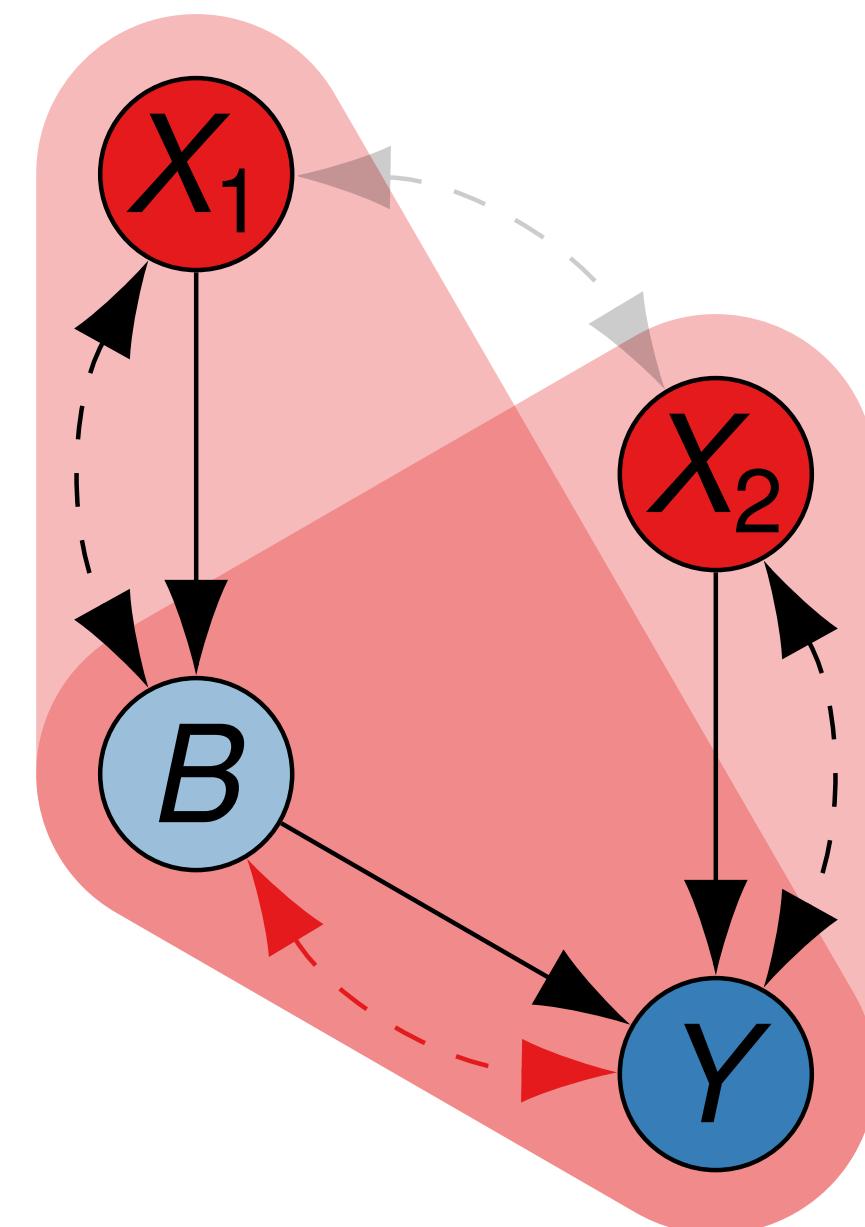


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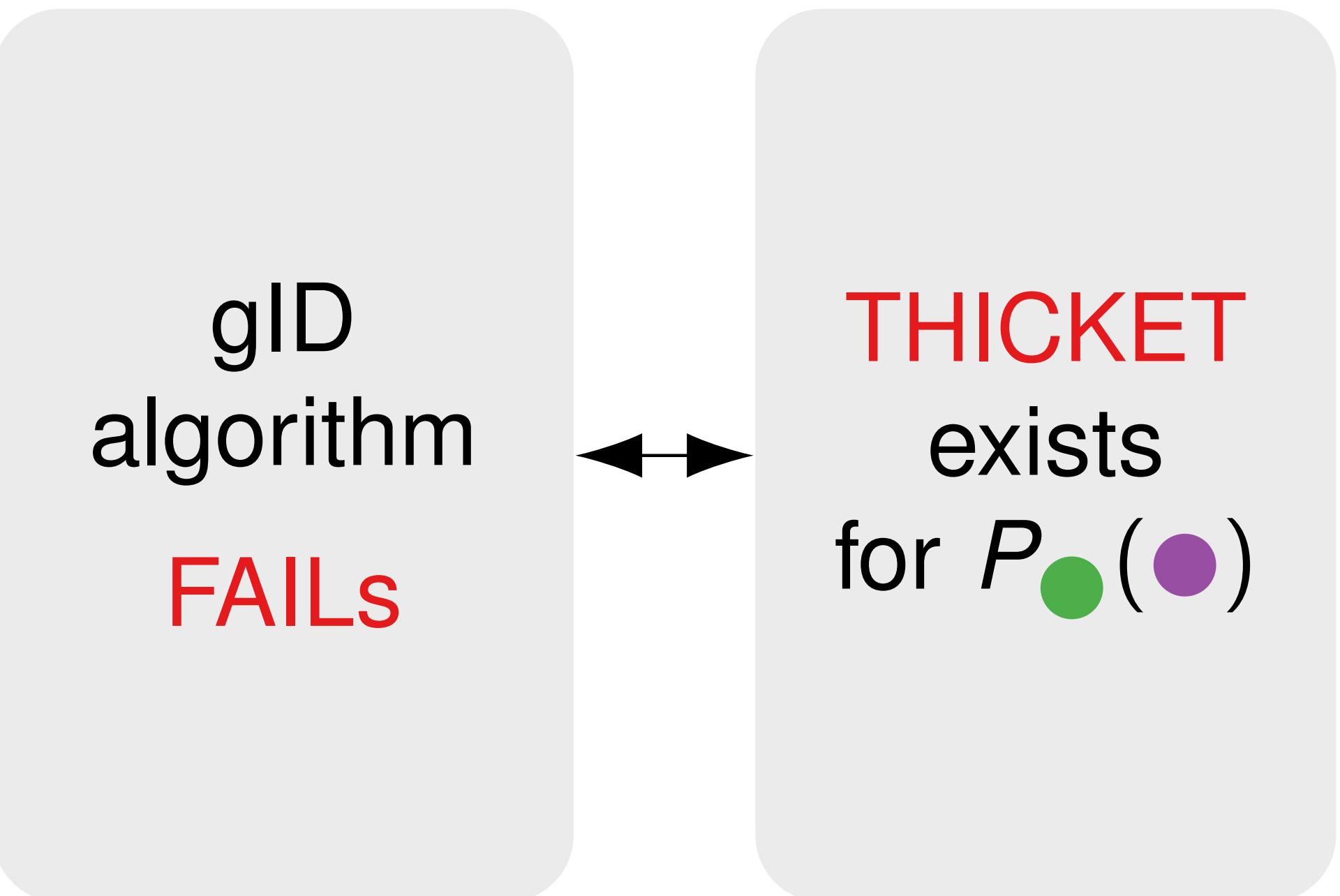
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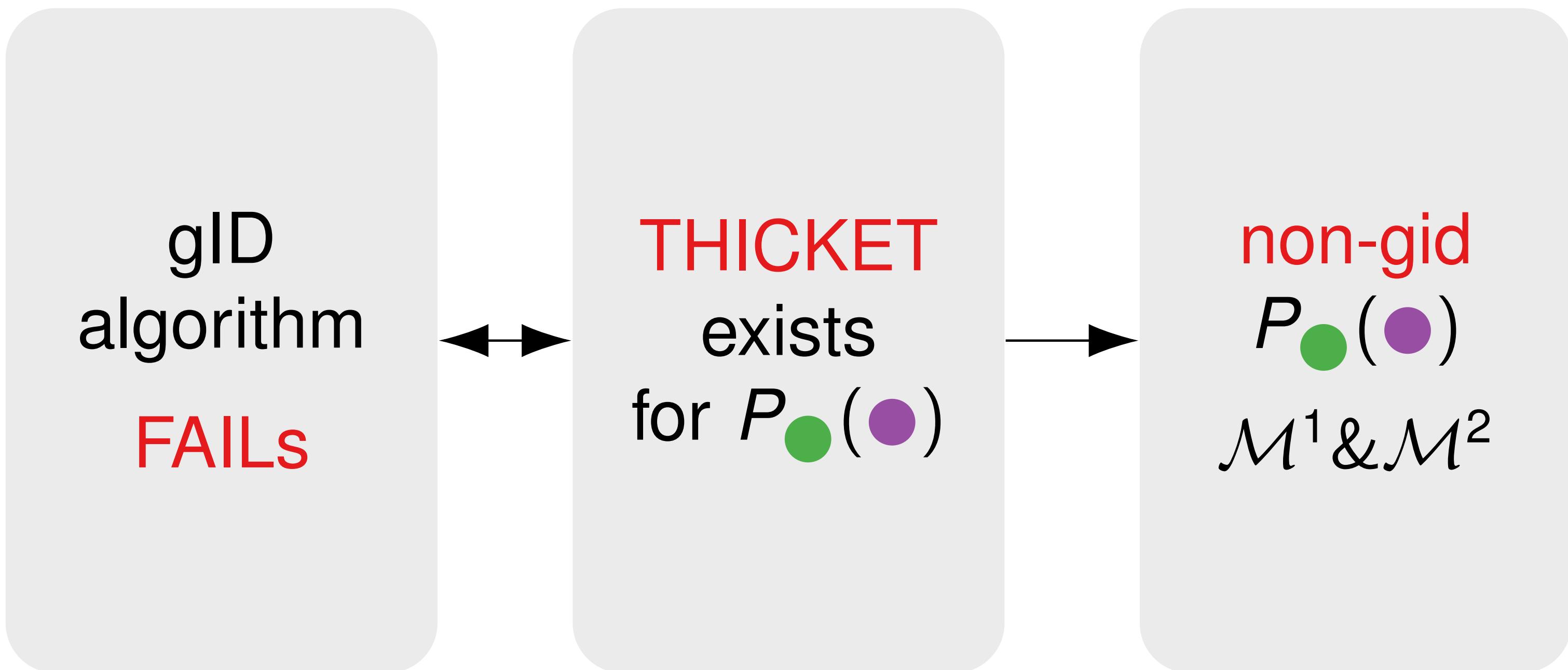
gID  
algorithm

FAILs

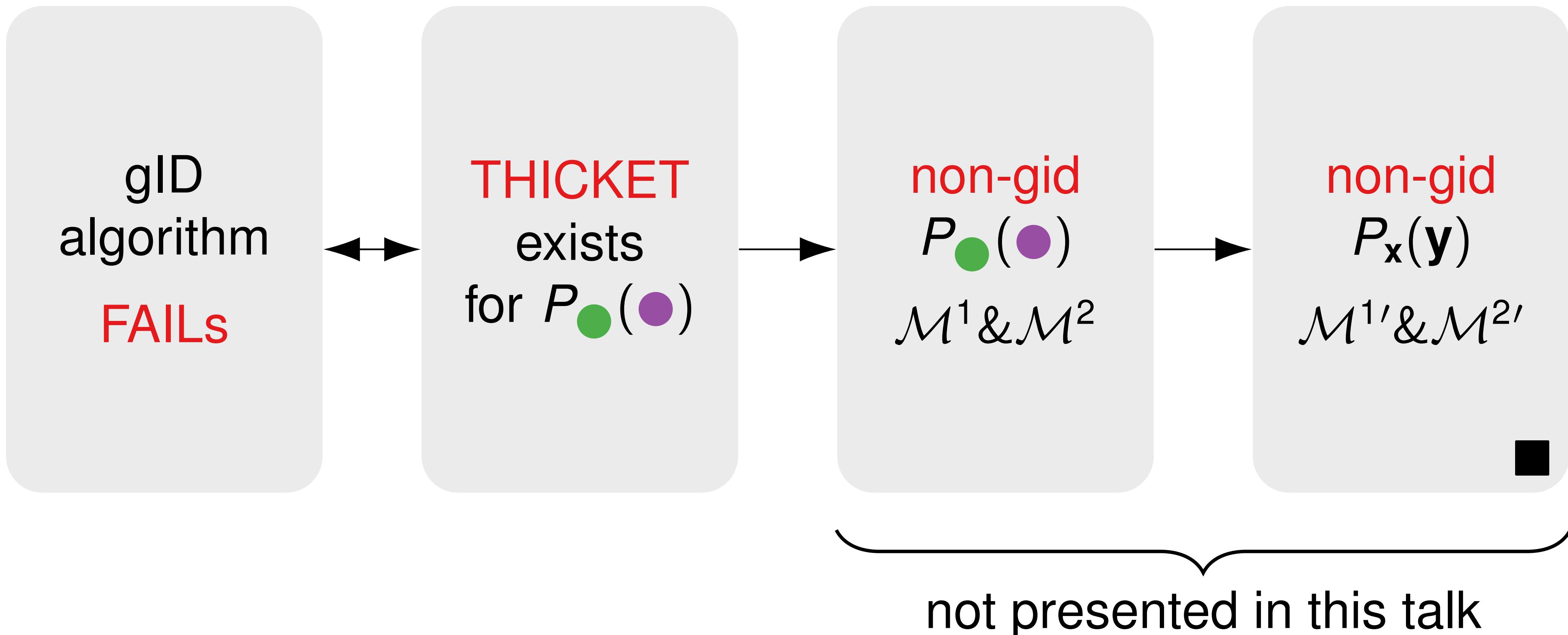
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# Conclusions

- We studied **general-identifiability** — causal effect identifiability given a causal graph and an **arbitrary combination of observational and experimental distributions**.
- ✓ a **necessary and sufficient** graphical **condition** ( $\exists$ thicket?).  
✓ a **sound and complete** **algorithm**
- Research Directions: finite-sample efficient formula, studying bounds for the causal effect when not-g-identifiable, incorporating functional assumptions, without a causal graph or partially-specified graphs.
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