

# Triangular Monotonic Generative Models Can Perform Causal Discovery

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## Triangular Monotonic (TMI) Maps

$$T(x) = \begin{bmatrix} T_1(x_1) \\ T_2(x_1, x_2) \\ \vdots \\ T_d(x_1, \dots, x_d) \end{bmatrix}$$

$T_1(x_1)$  ← Monotone  
 $T_2(x_1, x_2)$  ← Monotone in  $x_2$ ,  
 for all  $x_1$

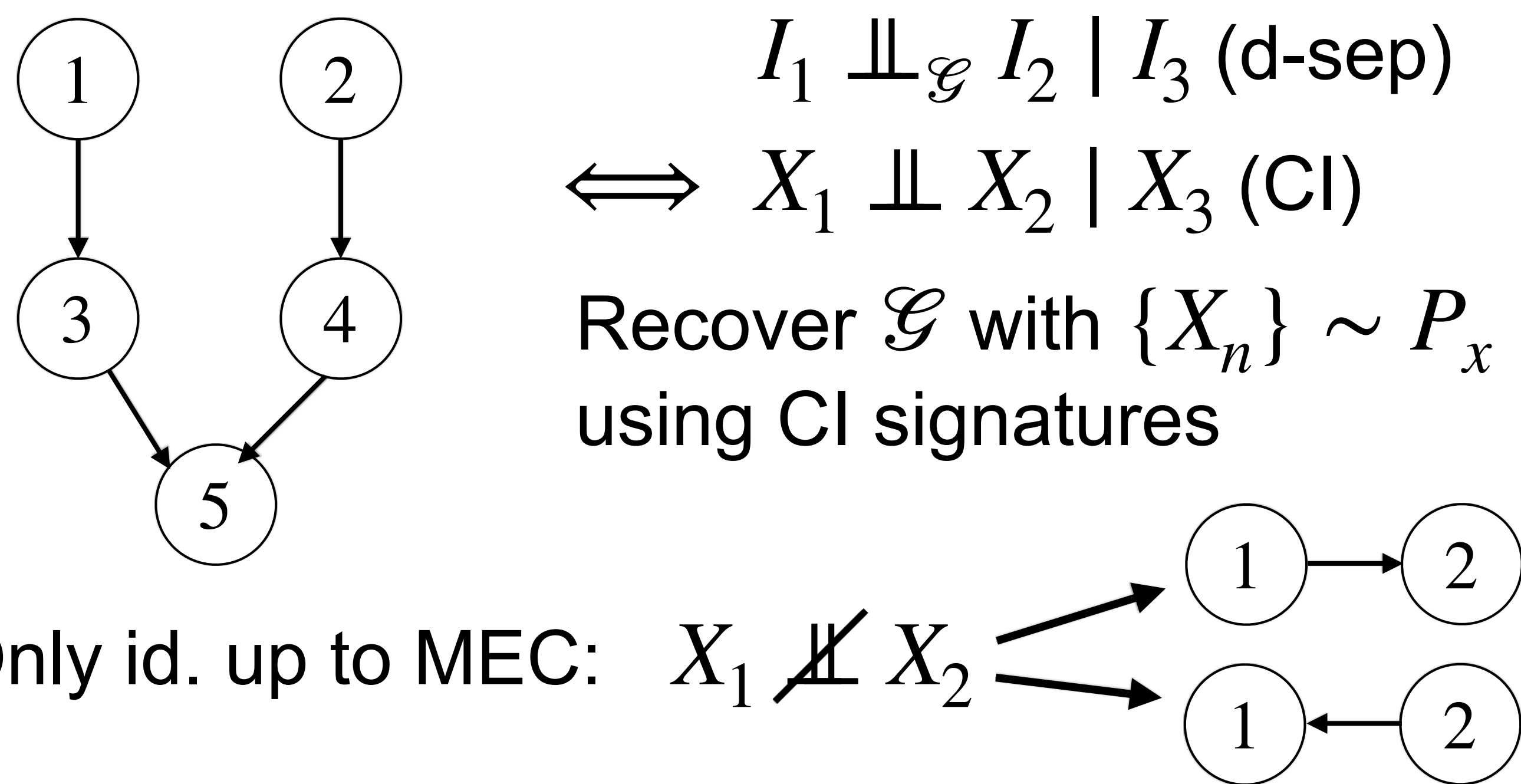
$P_\epsilon = \mathcal{U}[0,1]^d$      $T_{\#}P_x = P_\epsilon$ , full support ★  
 $T_i$  is the cond. CDF  $T_i(x_{\leq i}) = F_{X_i|X_{<i}}(x_i | x_{<i})$   
 Sparsity of  $T_i$  implies conditional independence

$$F_{X_i|X_{<i}} = F_{X_i|X_{<i} \setminus X_j} \iff X_i \perp\!\!\!\perp X_j | X_{<i} \setminus X_j \quad \star$$

Still holds if  $P_\epsilon$  is merely independent

## CI-based Structure Learning

Given DAG  $\mathcal{G}$  and faithful  $P_x$ :



Use a causal generative model (SCM) for  $P_x$ :

$$x_i = f(x_{pa(i)}, \epsilon_i), \text{ parents from } \mathcal{G}$$

Asymmetries in  $f$  can refine the MEC

## TMI Maps for Causal Discovery

TMI maps combine structure learning + SCM!

1) Permuting TMI maps can identify the MEC:

### Theorem (Raskutti + Uhler 2018):

The sparsest permutation of  $X$ , in terms of the number of CI's discovered in the sequence ★, identifies the MEC of  $\mathcal{G}$ .

I.e., finding  $\pi$  with the sparsest Jacobian gives adjacency matrix  $J_{T,\pi} \stackrel{Markov}{\sim} \mathcal{G}$

**Novelty:** don't need to assume an underlying SCM to do structure learning with Jacobian

2) TMIs can also fit SCMs:

If  $x_i = f(x_{pa(i)}, \epsilon_i)$  monotone in  $\epsilon_i$ , can write

$$(x_1, \dots, x_d) = T(\epsilon_1, \dots, \epsilon_d) \quad (\text{in practice, fit the abductive } T^{-1}, \text{ which is also TMI})$$

TMIs infer latent variables (★) if order correct

$$1 \rightarrow 2 : \hat{T}(x_1, x_2) = T^{-1}(x_1, x_2) = (\epsilon_1, \epsilon_2) \implies X_1 \perp\!\!\!\perp \epsilon_2$$

Incorrect order refines the MEC:

$$1 \leftarrow 2 : \text{true SCM not in model class!}$$

$$\hat{T}(x_1, x_2) = \begin{bmatrix} \hat{T}_1(x_1) \\ \hat{T}_2(x_1, x_2) \end{bmatrix} = (\hat{\epsilon}_1, \hat{\epsilon}_2) \implies X_1 \not\perp\!\!\!\perp \hat{\epsilon}_2$$

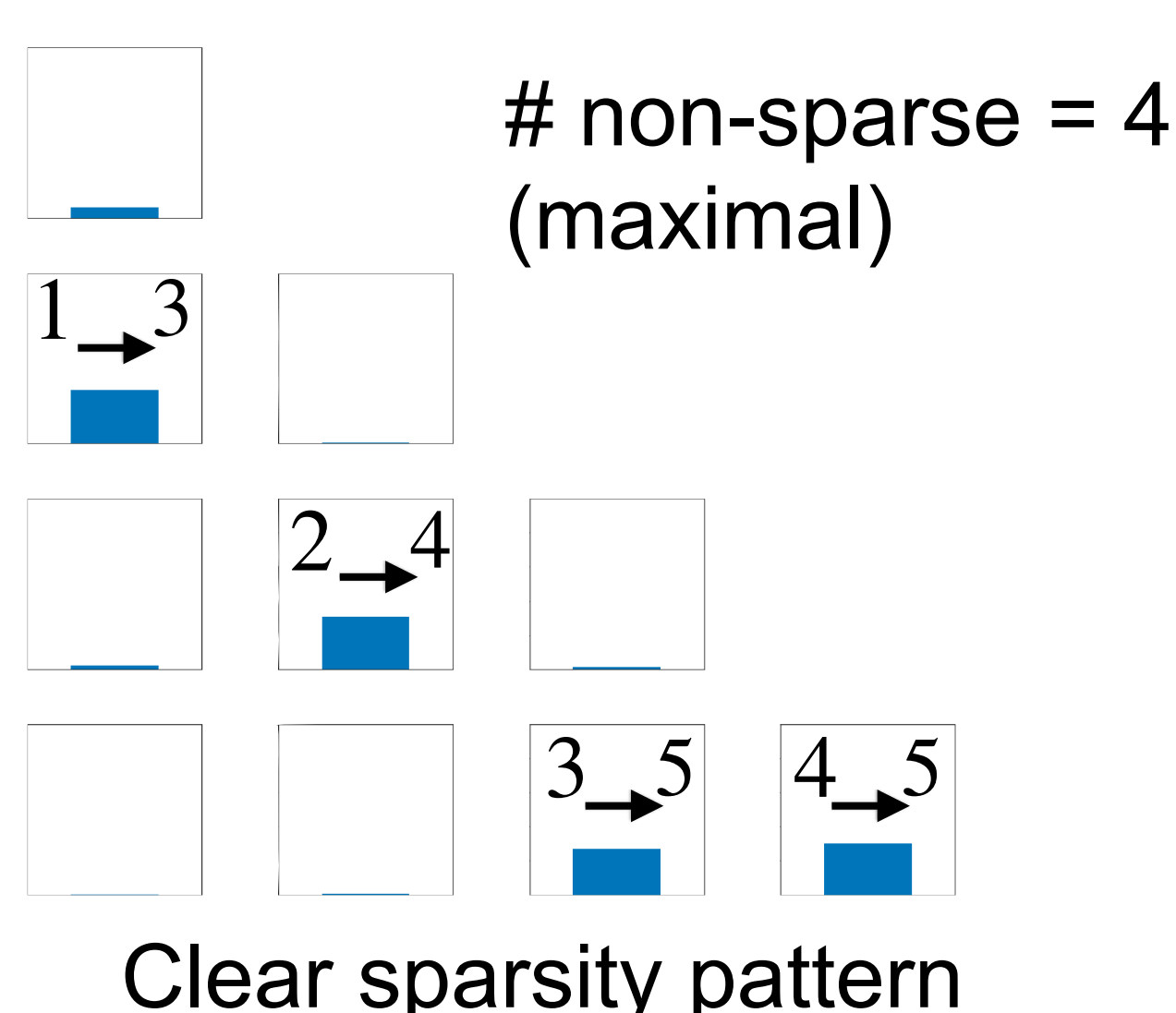
**Novelty:** TMI SCMs are less restrictive than the additive noise models typically used for CD

## Computational Challenges Ahead

Hard computational problems:

- Search over permutations 🤔
- How to threshold for sparsity 🤔
- Doing it for each permutation 🤔

Mean Jacobian for true  $\pi^*$



Mean Jacobian for incorrect  $\pi_r$

