

R.S. was supported by the Wellcome Trust [222836/Z/21/Z]. B.G. acknowledges the support of the UKRI AI programme, and the Engineering and Physical Sciences Research Council, for CHAI - EPSRC Causality in Healthcare AI Hub (grant no. EP/Y028856/1). A.K. is supported by UKRI (grant number EP/S023356/1), as part of the UKRI Centre for Doctoral Training in Safe and Trusted Al. A.D. acknowledges the support of G-Research's March 2025 grant.



Bayesian model selection allows for less restrictive functional assumptions when doing multivariate causal discovery on observational data

Bayesian Model Selection for Multivariate Causal Discovery

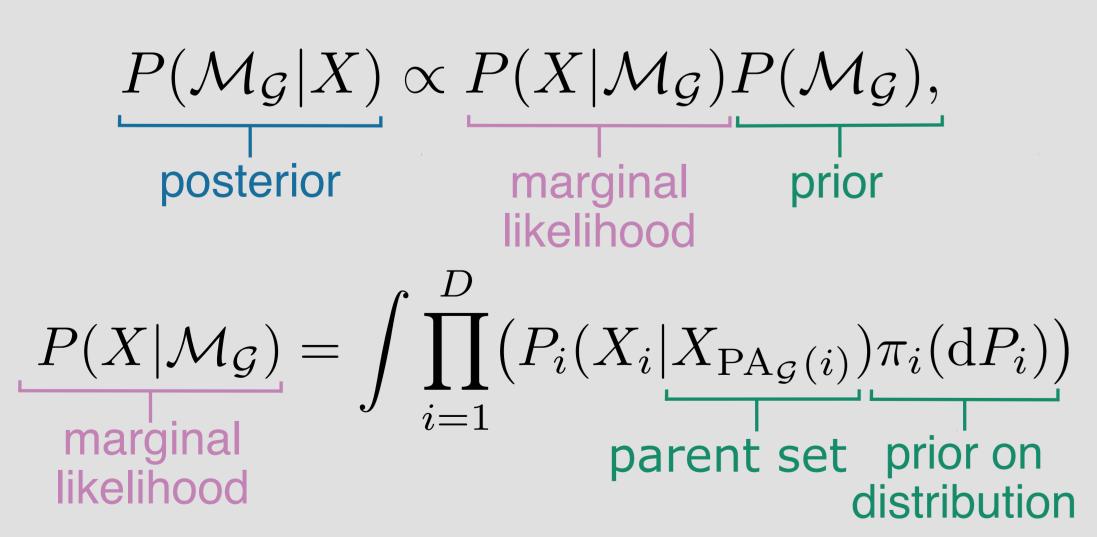
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Introduction

- Current causal discovery approaches require restrictive model assumptions to ensure identifiability, e.g. additive noise (ANM). In real-world data these restrictive assumptions are commonly violated, losing identifiability guarantees.
- Bayesian model selection has been proven to improve performance in the bivariate case by allowing more flexible functional assumptions.
- However, the naive discrete Bayesian model selection approach isn't feasible for the multivariate case where the number of possible graphs scales super-exponentially.
- We propose a continuous Bayesian model selection approach that scales well to large numbers of variables while still allowing more flexible functional assumptions.

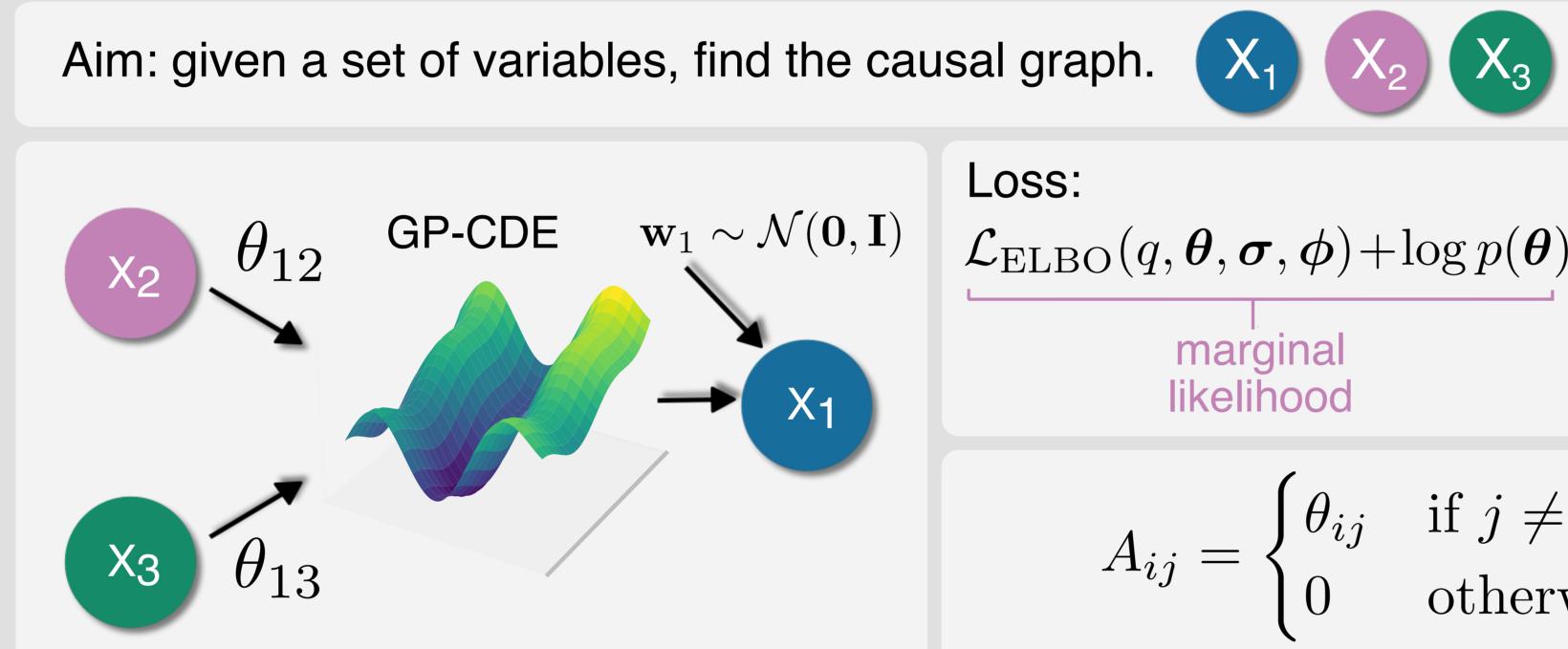
Bayesian model selection solves this problem



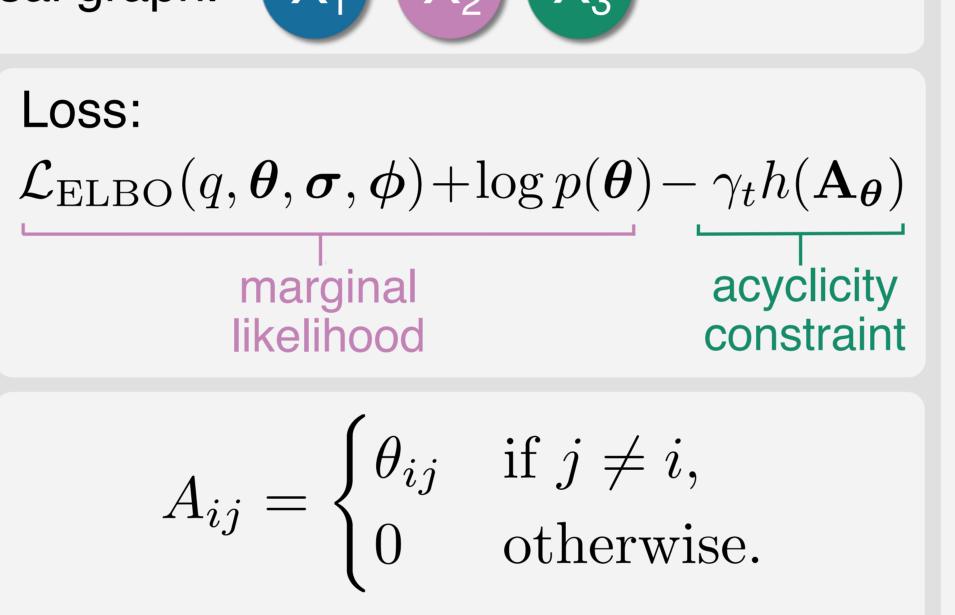
Causal model: $\mathcal{M}_{\mathcal{G}}$ Data: $X \in \mathbb{R}^D$

- Factorised prior on distributions encodes the independent mechanism (ICM) assumption.
- The marginal likelihood will prefer models whose ICM assumption aligns with the properties of the data generating process.

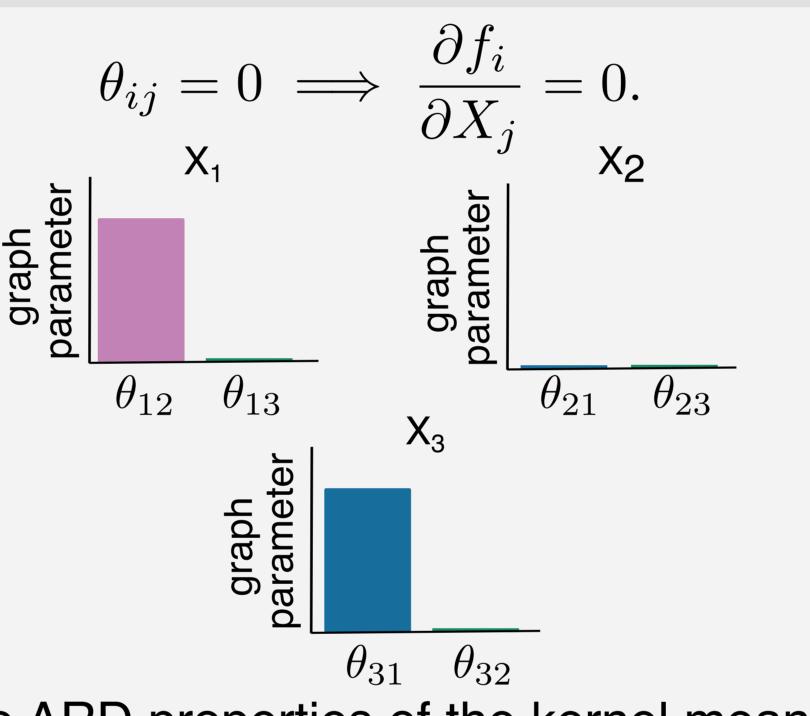
Causal Gaussian Process Conditional Estimator (CGP-CDE)



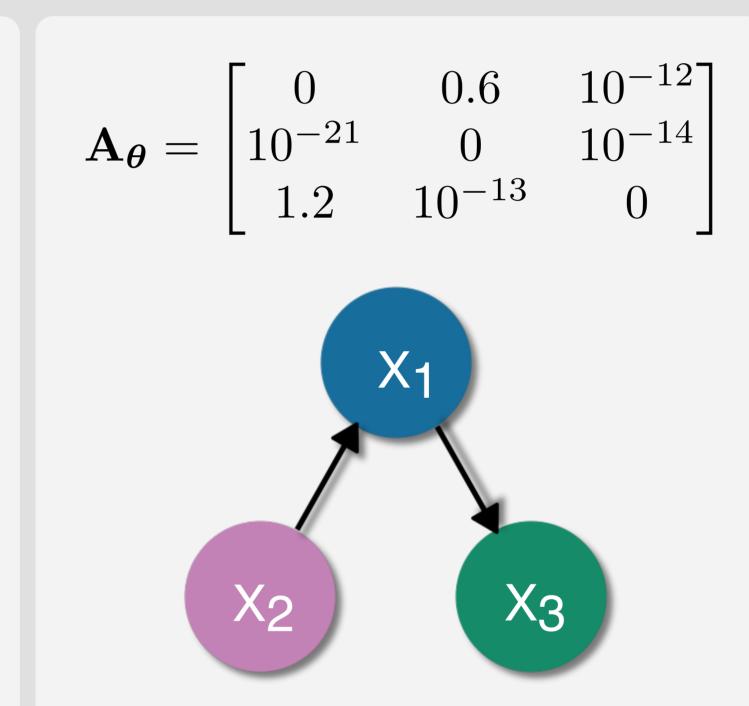
Fit a Gaussian process conditional estimator (GP-CDE) for each variable. Latent variable w₁ allows non-Gaussian and heteroskedastic densities.



An adjacency matrix is constructed from the graph parameters (inverse lengthscales).



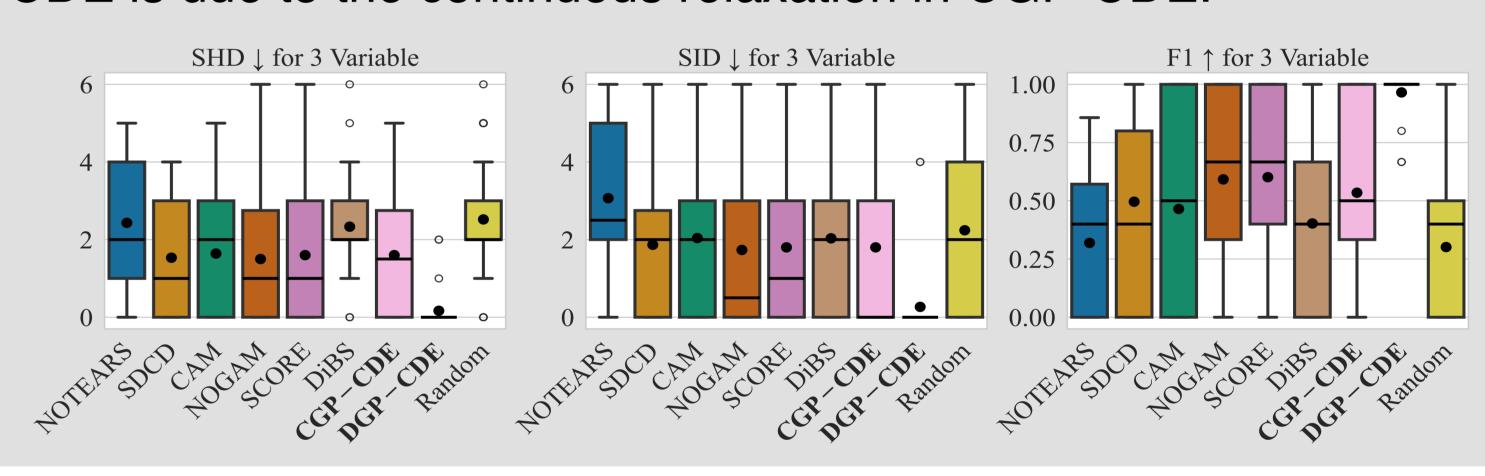
The ARD properties of the kernel mean edges that aren't evidenced by the data are "switched off" by making the graph parameter small.



The causal graph is extracted from the graph parameters via thresholding.

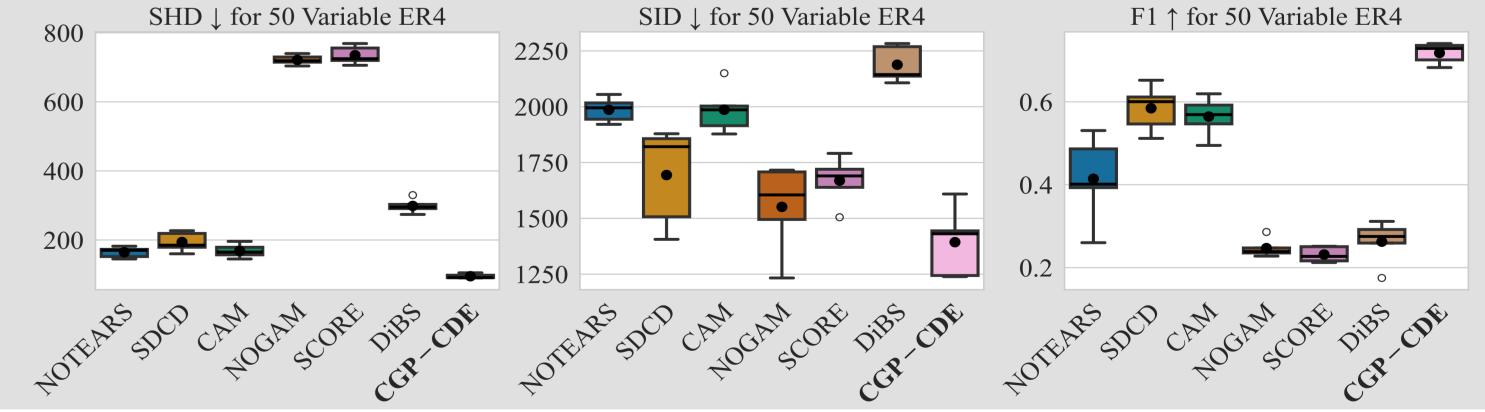
How bad is the continuous approximation?

- Excellent performance of discrete Bayesian model selection (DGP-CDE) shows the probability of error is small.
- The difference in performance between CGP-CDE and DGP-CDE is due to the continuous relaxation in CGP-CDE.



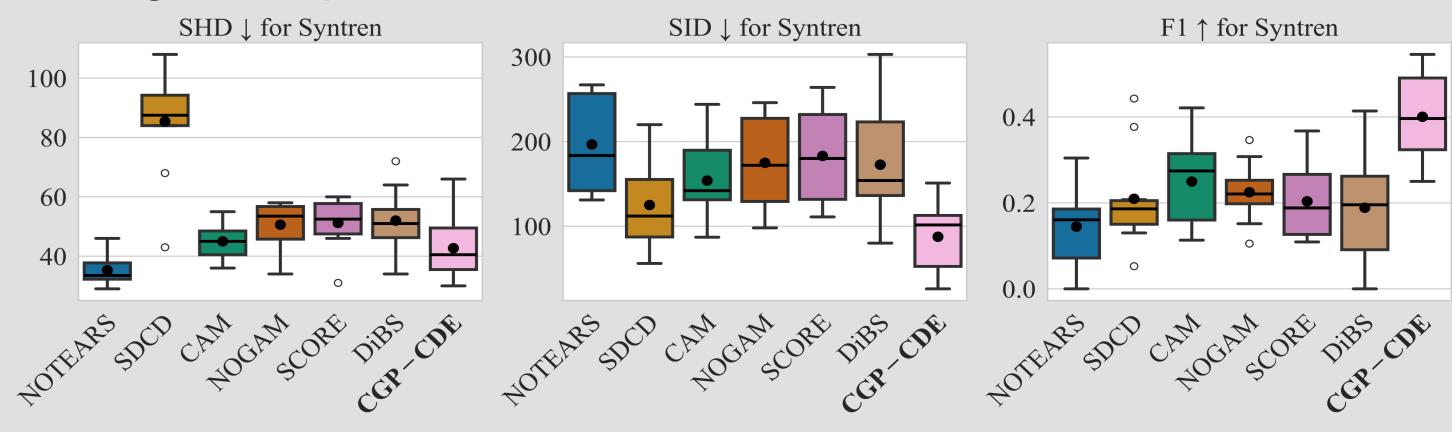
Does it scale to large numbers of variables?

• Experiments on ER graphs with 4 expected edges per variable and 50 variables. The CGP-CDE outperforms the other models.

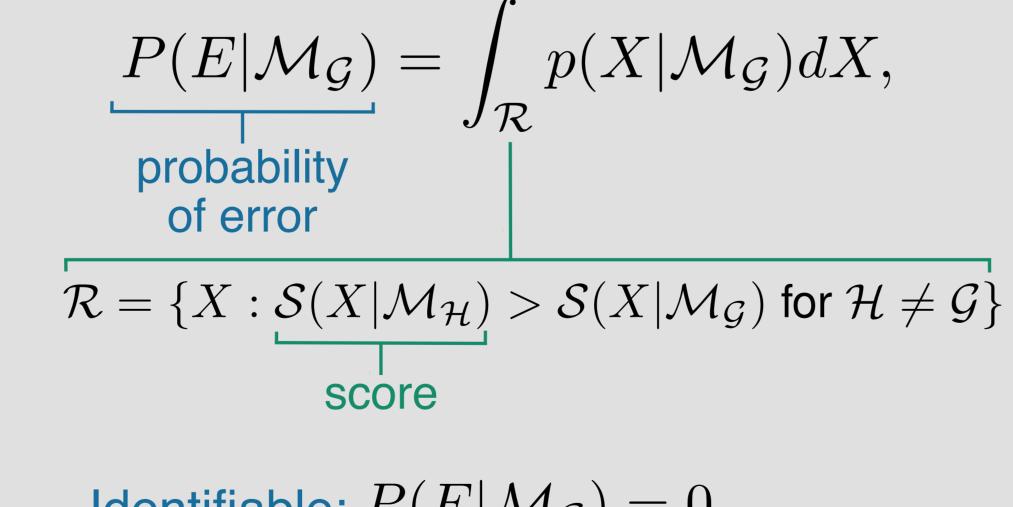


How well does it perform on semi-synthetic data?

• Experiments on Syntren dataset, derived from a gene regulatory network simulator which has 10 datasets of 20 nodes. The CGP-CDE again outperforms the other models.



Probability of error

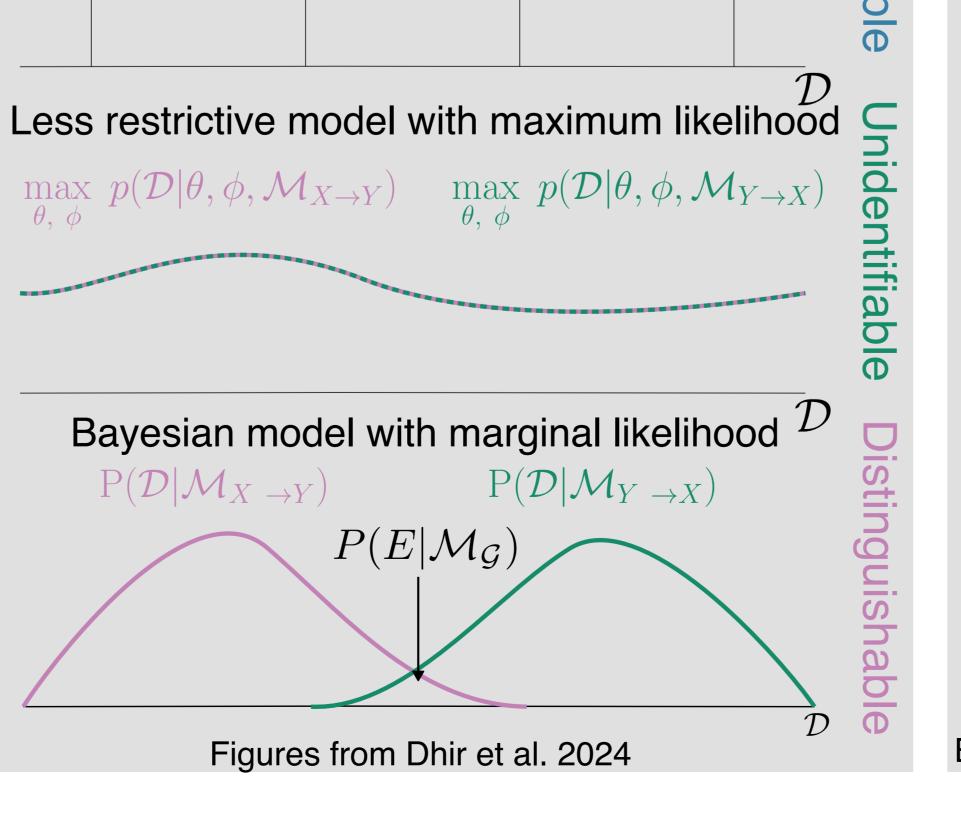


Identifiable: $P(E|\mathcal{M}_{\mathcal{G}})=0$

Unidentifiable: $P(E|\mathcal{M}_{\mathcal{G}}) = P(E|U(\mathcal{G}))$

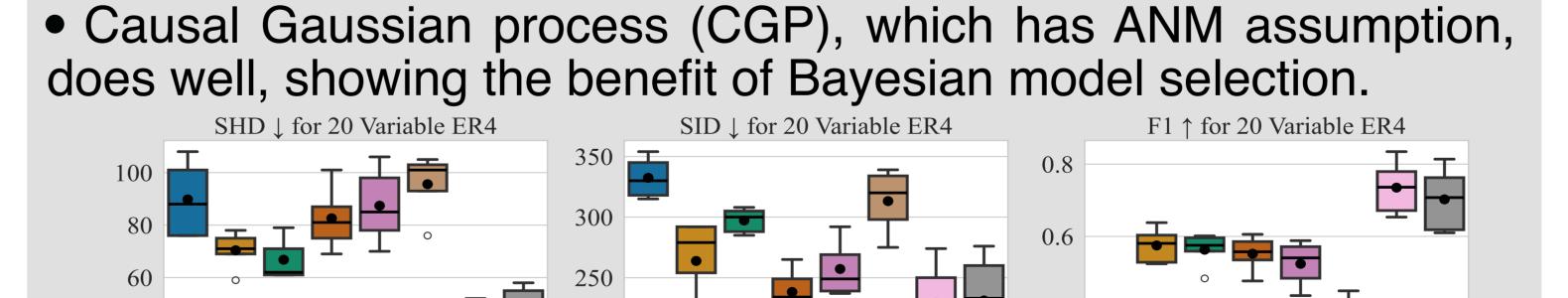
Distinguishable: $0 < P(E|\mathcal{M}_{\mathcal{G}}) < P(E|U(\mathcal{G}))$

uniformly random selected graph



ANM model with maximum likelihood

 $\max_{ heta,\ \phi}\ p(\mathcal{D}| heta,\phi,\mathcal{M}_{X o Y}') \qquad \max_{ heta,\ \phi}\ p(\mathcal{D}| heta,\phi,\mathcal{M}_{Y o X}') \qquad \mathbf{f Q}$



CGP-CDE outperforms ANM methods on ANM and non-ANM

Is performance on identifiable data worse?

