

# DAG-GNN: DAG Structure Learning with Graph Neural Networks

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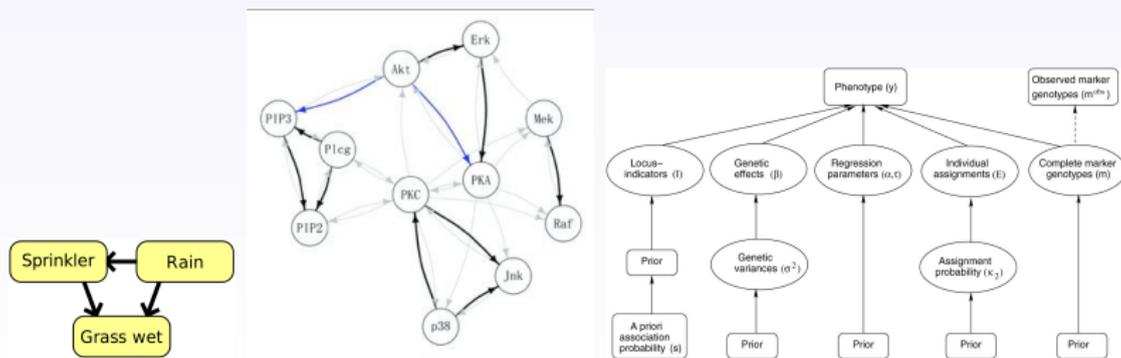
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# Motivation

The DAG learning problem is a vital part in causal inference:

- Let  $A \in \mathbb{R}^{m \times m}$  be the unknown weighted adjacency matrix of a DAG with  $m$  nodes.
- Given  $n$  identically distributed (i.i.d.) samples  $X^k \in \mathbb{R}^{m \times d}$ , from a distribution corresponding to  $A$ .
- Our focus is **to recovery the directed acyclic graph (DAG)  $A$  from  $X = \{X^1, \dots, X^n\}$ .**

However, DAG learning is proven to be NP-hard.



# Motivation

Conventional DAG learning methods:

- Perform score-and-search for discrete variables: with a constraint stating that the graph must be acyclic.
- Make a parametric (e.g. Gaussian) assumption for continuous variables: may result in model misspecification.

An equivalent acyclicity constraint was proposed by Zheng et al<sup>1</sup> (**NOTEARS**) for linear Structural Equation Model (SEM), by imposing a continuous penalty function

$$h(A) = \text{tr}(\exp(A \circ A)) - m.$$

We followed the framework of [1] to **formulate the problem as a continuous optimization**, with the following major contributions:

- 1 We developed a **deep generative model (VAE) parameterized by a novel graph neural network architecture (DAG-GNN)**.
- 2 We proposed an **alternative constraint  $h(A)$** .
- 3 The model is capable to capture **complex distributions** of data and to sample from them, and **naturally handles various data types**.

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<sup>1</sup>Zheng, X., Aragam, B., Ravikumar, P. K., & Xing, E. P. (2018). DAGs with NO TEARS: Continuous Optimization for Structure Learning. In Advances in Neural Information Processing Systems (pp. 9472-9483).

# Model Learning with Variational Autoencoder (VAE)

Our method learns the weighted adjacency matrix  $A$  of a DAG by using a deep generative model through maximizing the evidence lower bound (ELBO)

$$L_{\text{ELBO}} = \frac{1}{n} \sum_{k=1}^n L_{\text{ELBO}}^k,$$

$$L_{\text{ELBO}}^k \equiv -D_{\text{KL}}(q(Z|X^k) || p(Z)) + E_{q(Z|X^k)}[\log p(X^k|Z)].$$

The ELBO lends itself to a VAE: given  $X^k$ , the **encoder (inference model)** encodes it into a latent variable  $Z$  with density  $q(Z|X^k)$ ; and the **decoder (generative model)** reconstructs  $X^k$  from  $Z$  with density  $p(X^k|Z)$ .

Inspired by the linear SEM model

$$X = A^T X + Z, \text{ or, equivalently, } X = (I - A^T)^{-1} Z,$$

we propose a new graph neural network architecture for the decoder

$$\hat{X} = f_2((I - A^T)^{-1} f_1(Z)),$$

and the corresponding encoder

$$Z = f_4((I - A^T) f_3(X)).$$

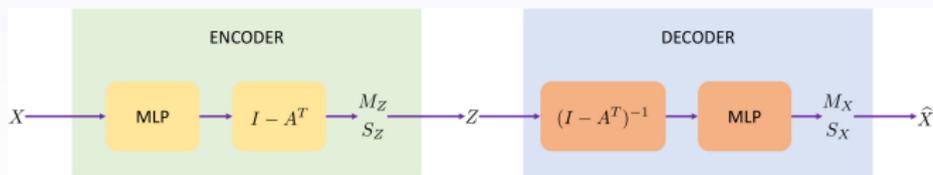
# Graph Neural Network (GNN) Architecture

For the **inference model (encoder)**  $Z = f_4((I - A^T)f_3(X))$ : we let  $f_3$  be a multilayer perceptron (MLP) and  $f_4$  be the identity mapping. Then the variational posterior  $q(Z|X)$  is a factored Gaussian with mean  $M_Z$  and standard deviation  $S_Z$ :

$$[M_Z | \log S_Z] = (I - A^T) \text{MLP}(X, W^1, W^2) := (I - A^T) \text{ReLU}(XW^1)W^2.$$

For the **generative model (decoder)**  $\hat{X} = f_2((I - A^T)^{-1}f_1(Z))$ : we let  $f_1$  be the identity mapping and  $f_2$  be an MLP. Then the likelihood  $p(X|Z)$  is a factored Gaussian with mean  $M_X$  and standard deviation  $S_X$ :

$$[M_X | \log S_X] = \text{MLP}((I - A^T)^{-1}Z, W^3, W^4) := \text{ReLU}((I - A^T)^{-1}ZW^3)W^4.$$



# A Robust Acyclicity Constraint

To further guarantee that the learnt  $A$  is acyclic, we propose an (alternative) equality constraint when maximizing the ELBO.

**Theorem:** Let  $A \in \mathbb{R}^{m \times m}$  be the (possibly negatively) weighted adjacency matrix of a directed graph. For any  $\alpha > 0$ , the graph is acyclic if and only if

$$h(A) = \text{tr}[(I + \alpha A \circ A)^m] - m = 0.$$

Here  $\alpha$  may be treated as a hyperparameter.

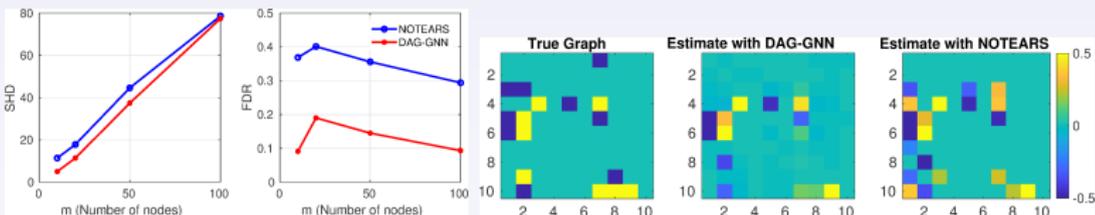
When the eigenvalues of  $A \circ A$  have a large magnitude, by taking sufficiently small constant  $\alpha$ ,  $(I + \alpha A \circ A)^m$  is more stable than  $\exp(A \circ A)$ :

**Theorem:** Let  $\alpha = c/m > 0$  for some  $c$ . Then for any complex  $\lambda$ , we have  $(1 + \alpha|\lambda|)^m \leq e^{c|\lambda|}$ .

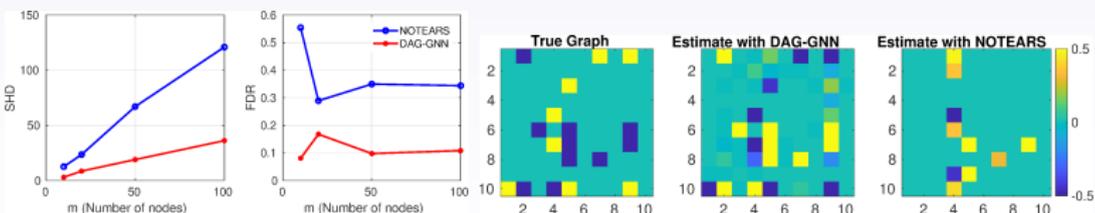
In practice,  $\alpha$  depends on  $m$  and an estimation of the largest eigenvalue of  $A \circ A$  in magnitude.

# Nonlinear and vector value datasets

- Nonlinear synthetic data:** generated by  $X = A^T \cos(X + \mathbf{1}) + Z$ :



- Vector value data**  $X^k \in \mathbb{R}^{m \times d}$ ,  $d > 1$ : generated by  $\tilde{x} = A^T \tilde{x} + \tilde{z}$ ,  $x^k = u^k \tilde{x} + v^k + z^k$  and  $X = [x^1 | x^2 | \dots | x^d]$ :



## Discrete value datasets

The proposed model naturally handles **discrete variables**. Assuming that each variable has a finite support of cardinality  $d$ , let  $p(X|Z)$  be a factored categorical distribution with probability matrix  $P_X$ , one embedding layer is added to the encoder and the decoder is modified as:

$$P_X = \text{softmax}(\text{MLP}((I - A^T)^{-1}Z, W^3, W^4)).$$

The solver is compared with the state-of-the-art exact DAG solver GOPNILP<sup>2</sup> on 3 benchmark datasets:

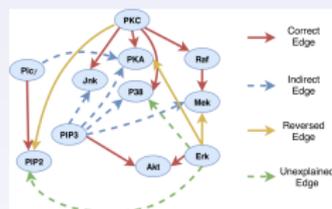
Dataset	$m$	Groundtruth	GOPNILP	DAG-GNN
Child	20	-1.27e+4	-1.27e+4	-1.38e+4
Alarm	37	-1.07e+4	-1.12e+4	-1.28e+4
Pigs	441	-3.48e+5	-3.50e+5	-3.69e+5

Table : BIC scores on benchmark datasets of discrete variables.

<sup>2</sup>Cussens, J., Haws, D., & Studeny, M. (2017). Polyhedral aspects of score equivalence in Bayesian network structure learning. *Mathematical Programming*, 164(1-2), 285-324.

Applied to a **bioinformatics dataset**<sup>3</sup> for the discovery of a protein signaling network:

Method	SHD	# Predicted edges
FGS	22	17
NOTEARS	22	16
DAG-GNN	19	18



Applied to a **knowledge base (KB) schema dataset**<sup>4</sup>. The nodes of which are relations and the edges indicate whether one relation suggests another.

film/ProducedBy	⇒	film/Country
film/ProductionCompanies	⇒	film/Country
person/Nationality	⇒	person/Languages
person/PlaceOfBirth	⇒	person/Languages
person/PlaceOfBirth	⇒	person/Nationality
person/PlaceLivedLocation	⇒	person/Nationality

<sup>3</sup>Sachs, K., Perez, O., Pe'er, D., Lauffenburger, D. A., & Nolan, G. P. (2005). Causal protein-signaling networks derived from multiparameter single-cell data. *Science*, 308(5721), 523-529.

<sup>4</sup>Toutanova, K., Chen, D., Pantel, P., Poon, H., Choudhury, P., & Gamon, M. (2015). Representing text for joint embedding of text and knowledge bases. In *Proceedings of the 2015 Conference on Empirical Methods in Natural Language Processing* (pp. 1499-1509).

# Thank you for your attention.

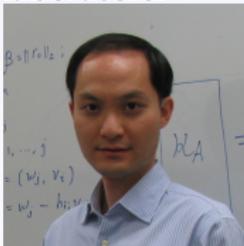
The code is available at <https://github.com/fishmoon1234/DAG-GNN>.

For further details and questions, please come to our poster session:

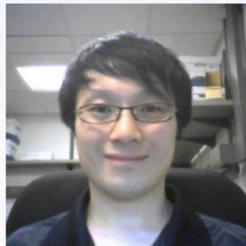
**This evening 06:30 – 09:00 PM, Pacific Ballroom #215.**

## Acknowledgement

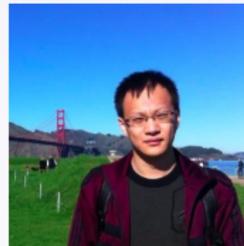
- **Collaborators:**



Jie Chen



Tian Gao



Mo Yu

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