# DAGs with NO TEARS: Continuous Optimization for Structure Learning 

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## Graphical Models

- Graphical Models are families of multivariate distributions with compact representations, scaling to very large numbers of variables
- Why do we need compact representations?
- even for binary random variables, specification of a general multivariate distribution over $p$ variables requires $O\left(2^{p}\right)$ values
- In general, parametrizing higher-order dependencies among random variables scales poorly (typically exponentially) with number of variables
- Graphical models only require "local" specifications based on graph neighborhoods (associating variables to graph nodes), and hence scale well to large number of variables
- Undirected graphical models, also called Markov networks, or Markov random fields: family represented by an undirected graph
- Directed graphical models, also called Bayesian networks: family represented by a directed acyclic graph (DAG)


## Directed Graphical Models

$X=\left(X_{1}, \ldots, X_{p}\right) \sim$ directed graphical model with DAG $G=(V, E)$ if:

$$
P(X ; G)=\prod_{j=1}^{p} P\left(X_{j} \mid X_{\mathrm{pa}_{j}}\right),
$$

- $\mathrm{pa}_{j}$ is the set of parents of node $j \in V$
- associating variables $X_{j}$, for $j \in[p]$, with nodes $j \in V$
- Only requires local specifications of conditional distributions of variables given its "parent" variables


## Directed Graphical Models

The DAG $G$ encodes the conditional independence assumptions satisfied by resulting distributions simply as:

$$
X_{j} \Perp X_{\mathrm{nd}_{j}} \mid X_{\mathrm{pa}_{j}}, \forall j \in V
$$

- $\mathrm{pa}_{j}$ is set of parents of node $j \in V$
- $\mathrm{nd}_{j}$ is set of non-descendants of node $j \in V$
- Edges connote "direct dependence" that is more meaningful than high correlation; underlying DAG G an object of interest even when the full multivariate distribution is not
- Applications across biology (Sachs et al., 2005), genetics (Zhang et al., 2013), causal inference (Spirtes et al., 2000), artificial intelligence (Koller and Friedman, 2009), and many more


## Learning DAGs

- Graphical models: compact models of $p\left(x_{1}, \ldots, x_{d}\right)$

- Structure learning: what graph fits the data best?



## Learning DAGs

- Two main classes of approaches
- Conditional Independence Test based
- test which conditional independences hold in the data, find graph that best corresponds to these
- caveats: many more conditional independences might hold in data ("lack of faithfulness"), sensitive to failure of individual tests + multiplicity of tests; computationally less scalable
- Score based
- search for graph that optimizes some score (measuring goodness of fit of graph to data)
- typically local search/greedy algorithms that greedily build graph
- NP-hard in general, need many model specific heuristics to "get it to work"


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- caveats: many more conditional independences might hold in data ("lack of faithfulness"), sensitive to failure of individual tests + multiplicity of tests; computationally less scalable
- Score based
"The disadvantage of the score-based approaches (for Bayesian networks) is that they pose a search problem that may not have an elegant and efficient solution" .... Koller and Friedman (2009, pp. 785)
- NP-hard in general, need many model specific heuristics to "get it to work"


# Learning Bayesian Networks vs Markov Networks 

|  | Markov <br> Networks | Bayesian Networks |
| :---: | :---: | :---: |
| Cond. Indep. <br> test based | folklore | Spirtes and Glymour <br> $(1991)$ |
| Score based <br> (local search) | Pietra et al (1997) | Heckerman et al |
| (1995) |  |  |

## Learning DAGs: Problem Setup

- Bayesian network (BN) $G$ with $d$ nodes:

$$
p_{\text {joint }}\left(x_{1}, \ldots, x_{d} ; G\right)=\prod_{j=1}^{d} p_{\text {cond }}\left(x_{j} \mid \boldsymbol{x}_{p a(j)}\right)
$$

Structural Equation Models (SEMs) specify the form of these conditional distributions:

$$
\begin{aligned}
X_{j} & =T\left(X_{\mathrm{pa}_{j}}, \epsilon_{j}\right) \\
\epsilon_{j} & \sim \text { noise distribution }
\end{aligned}
$$

We will be considering SEMs parameterized by a weighted adjacency matrix $\mathbf{W}$

## Linear SEMs

- Weighted adjacency matrix $W \in \mathbb{R}^{d \times d}$ :

$$
W=\left[\begin{array}{ccccc}
w_{1 \rightarrow 1} & \cdots & w_{1 \rightarrow j} & \cdots & w_{1 \rightarrow d} \\
\vdots & \ddots & \vdots & \ddots & \vdots \\
w_{d \rightarrow 1} & \cdots & w_{d \rightarrow j} & \cdots & w_{d \rightarrow d}
\end{array}\right]
$$

The $j$-th column $w_{j}$ : edge weights from $p a(j)$ to $j$.

- Linear BN:

$$
x_{j}=T\left(\boldsymbol{x}_{p a(j)}, \varepsilon_{j}\right)=\underbrace{\boldsymbol{x}^{\top} w_{j}}_{\text {linear }}+\underbrace{\varepsilon_{j}}_{\text {zero mean }}
$$

## Generalized Linear SEMs

- Weighted adjacency matrix $W \in \mathbb{R}^{d \times d}$ :

$$
W=\left[\begin{array}{ccccc}
w_{1 \rightarrow 1} & \cdots & w_{1 \rightarrow j} & \cdots & w_{1 \rightarrow d} \\
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\end{array}\right]
$$

The $j$-th column $w_{j}$ : edge weights from $\mathrm{pa}(j)$ to $j$.

- Generalized Linear BN:

$$
P\left(x_{j} \mid \boldsymbol{x}_{p a(j)}\right)=h\left(x_{j}\right) \exp \left(g\left(x_{j}\right)\left(\boldsymbol{x}^{\top} w_{j}\right)-A\left(\boldsymbol{x}^{\top} w_{j}\right)\right),
$$

so that:

$$
E\left(x_{j} \mid x_{p a(j)}\right)=T(\underbrace{\boldsymbol{x}^{\top} w_{j}}_{\text {linear }}),
$$

for some function $T(\cdot)$. Examples: Logistic, Poisson, $\cdots$

## Generative Process: Linear SEMs

- Drawing one sample $\boldsymbol{x}$ :

For each node $j$ in topological order of $G$,

$$
\begin{aligned}
& \varepsilon_{j} \sim \text { noise distribution } \\
& x_{j}=T\left(\boldsymbol{x}_{p a(j)}, \varepsilon_{j}\right)
\end{aligned}
$$

- Let $X \in \mathbb{R}^{n \times d}$ be collection of $n$ such samples:

$$
X=\left[\begin{array}{ccc}
- & \boldsymbol{x}^{(1)} & - \\
& \vdots & \\
- & \boldsymbol{x}^{(n)} & -
\end{array}\right]
$$

## Generative Process: Linear SEMs

- For $j$-th dimension of $i$-th sample,

$$
x_{j}^{(i)}=\boldsymbol{x}^{(i)^{T}} \boldsymbol{w}_{j}+\varepsilon_{j}^{(i)}
$$

- Collecting all $i \in[n], j \in[d]$,

$$
\begin{gathered}
X=X W+E \\
(n \times d) \quad(n \times d)(d \times d)(n \times d)
\end{gathered}
$$

- Natural loss function:

$$
\ell(W ; X)=\frac{1}{2 n}\|X-X W\|_{F}^{2}
$$

## M-Estimation for DAGs

Given $X \in \mathbb{R}^{n \times d}$, solve

$$
\begin{aligned}
\min _{W \in \mathbb{R}^{d \times d}} & \ell(W ; X) \\
\text { s.t. } & G(W) \in D A G
\end{aligned}
$$

- Loss function: log-likelihood of data, with respect to SEM model, given weighted adjacency matrix W
- Constraint: that W correspond to some DAG G
- leads to difficult combinatorial optimization problem


## Smooth Characterization of DAGs?

## Smooth Characterization of DAG

Can we find a smooth function $h: \mathbb{R}^{d \times d} \rightarrow \mathbb{R}$ such that

$$
h(W)=0 \Longleftrightarrow G(W) \in D A G
$$

holds?

- will enable solvers based on continuous optimization, similar to Markov networks, in contrast to solving via constraint estimation, or local search in space of DAGs


## Finite Power Series?

Consider binary adjacency matrix $B \in\{0,1\}^{d \times d}$

- Idea:

$$
\left(B^{k}\right)_{i j}=\text { num of } k \text {-step paths from } i \text { to } j
$$

In other words,

$$
\operatorname{tr}\left(B^{k}\right)=0 \Longleftrightarrow \text { no } k \text {-cycles }
$$

- Candidate:

$$
h(B)=\operatorname{tr}\left(\sum_{k=1}^{d} B^{k}\right)=0 \Longleftrightarrow G(B) \in D A G
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$$

Caveat: number of $\mathbf{k}$ cycles increases exponentially (ill conditioned)

## Infinite Power Series?

- Idea: push $k$ to infinity

$$
\sum_{k=0}^{\infty} B^{k}=(I-B)^{-1}
$$

- Candidate:

$$
h(B)=\operatorname{tr}(I-B)^{-1}-d=0 \Longleftrightarrow G(B) \in D A G
$$

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$$

Caveat: requires invertibility of I-B (i.e. that spectral radius of $B<1$ )

## Matrix Exponential

- Idea: is there a series that always converges? Yes!

$$
e^{B}=I+B+\frac{1}{2!} B^{2}+\frac{1}{3!} B^{3}+\cdots
$$

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$$

Caveat: requires that adjacency matrix B be binary

## Matrix Exponential for General Adjacency Matrices

- Idea: for nonnegative matrix $S \in \mathbb{R}_{+}^{d \times d}$,

$$
\begin{aligned}
& \left(S^{k}\right)_{i j}=\text { sum of weight products long } k \text {-step paths from } i \text { to } j \\
& \qquad \operatorname{tr}\left(S^{k}\right)=0 \Longleftrightarrow \text { no } k \text {-cycles }
\end{aligned}
$$

- Real to nonnegative:

$$
S=W \circ W
$$

- Candidate:

$$
h(W)=\operatorname{tr}\left(e^{W \circ W}\right)-d=0 \Longleftrightarrow G(W) \in D A G
$$

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Can we find a smooth function $h: \mathbb{R}^{d \times d} \rightarrow \mathbb{R}$ such that

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holds?
Answer: Yes!

$$
h(W)=\operatorname{tr}\left(e^{W \circ W}\right)-d=0 \Longleftrightarrow G(W) \in D A G
$$

Furthermore, it has a simple gradient

$$
\nabla h(W)=\left(e^{W \circ W}\right)^{T} \circ 2 W
$$

## New M-estimator for DAGs

Given $X \in \mathbb{R}^{n \times d}$, solve

$$
\begin{aligned}
\min _{W \in \mathbb{R}^{d \times d}} & \ell(W ; X) \\
\text { s.t. } & h(W)=0
\end{aligned}
$$

## Optimization Algorithm: Augmented Lagrangian

- Solve an equivalent augmented form

$$
\begin{aligned}
\min _{W \in \mathbb{R}^{d \times d}} & \ell(W ; X)+\frac{\rho}{2} h^{2}(W) \\
\text { s.t. } & h(W)=0
\end{aligned}
$$

- Lagrangian:

$$
L(W, \alpha)=\ell(W ; X)+\frac{\rho}{2} h^{2}(W)+\alpha h(W)
$$

- Solve the dual:

$$
\min _{W} \max _{\alpha} L(W, \alpha)=\underbrace{\max _{\alpha} \overbrace{\min }^{\text {smooth, unconstrained }} L(W, \alpha)}_{\text {1d linear maximization }}
$$

## NO TEARS

## Algorithm 1 Augmented Lagrangian

- Input: $\ell, \nabla \ell, h, \nabla h$
- For $t=1,2,3, \ldots$.
- Solve primal $W_{t+1} \leftarrow \operatorname{argmin}_{W} L\left(W, \alpha_{t}\right)$.
- Dual ascent $\alpha_{t+1} \leftarrow \alpha_{t}+\rho h\left(W_{t+1}\right)$.

NOTEARS $=$ Non-combinatorial Optimization via Trace Exponential and Augmented lagRangian for Structure learning

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NOTEARS $=$ Non-combinatorial Optimization via Trace Exponential and Augmented lagRangian for Structure learning
(Corollary 11.2.1, Nemirovski, 1999) (loosely)
For $\rho$ large enough, and with starting point $\alpha_{0}$ near an optimum $a^{\star}$, the updates converge to $a^{*}$ linearly.

## NO TEARS

```
def notears_simple(x, max_iter=100, h_tol=1e-8, w_threshold=0.3):
    n, d = X.shape
    w_est, w_new = np.zeros(d * d), np.zeros(d * d)
    rho, alpha, h, h_new = 1.0, 0.0, np.inf, np.inf
    bnds = [(0, 0) if i == j else (None, None) for i in range(d) for j in range(d)]
    for _ in range(max_iter):
        while rho < 1e+20:
            sol = sopt.minimize(_func, w_est, method='L-BFGS-B', jac=_grad, bounds=bnds)
            w_new = sol.x
            h_new = _h(w_new)
            if h_new }\mp@subsup{}{}{\prime}>0.25 * h
            rho *= 10
            else:
                    break
        w_est, h = w_new, h_new
        alpha += rho * h
        if h <= h_tol:
            break
    w_est[np.abs(w_est) < w_threshold] = 0
    return w_est.reshape([d, d])
```

30 lines (function, gradient) +20 lines (optimize) $\approx 50$ lines in total
In contrast to 1000s of lines of combinatorial optimization code (with model-specific heuristics)

Code available at: https://github.com/xunzheng/notears

## Also amenable to structured sparsity constraints on $G$

Prefer sparse graphs:

$$
\begin{aligned}
\min _{W \in \mathbb{R}^{d \times d}} & \ell(W ; X)+\lambda\|W\|_{1} \\
\text { s.t. } & h(W)=0
\end{aligned}
$$

No longer smooth!
Proximal Quasi-Newton for augmented Lagrangian:

$$
\begin{aligned}
& \min _{W} L(W, \alpha) \\
= & \min _{W} \underbrace{\ell(W ; X)+\frac{\rho}{2} h^{2}(W)+\alpha h(W)}_{\text {smooth }}+\lambda\|W\|_{1}
\end{aligned}
$$

## Caveats with M-estimator

- Constraint set is non-convex
- Computing matrix exponential is $\mathrm{O}\left(\mathrm{d}^{3}\right)$
- With typical continuous optimization algorithms, constraints are only satisfied up to some tolerance
- due to which we add a post-processing thresholding step to weighted adjacency matrix


## Experiments

- Random graphs: Erdos-Renyi (ER), Scale Free (SF)
- Samples $\mathrm{n}=\{20,1000\}$, variables $\mathrm{d}=\{10,20,100,200\}$
- Baseline: FGS (Fast Greedy Search; Ramsey et al 2016); state of art implementation of greedy equivalent search (GES; Chickering 2008); has been known to outperform other local search techniques
- Metrics:
- False Discovery Rate (FDR): \#false edges/\#predicted edges
- Structural Hamming Distance (SHD): \#false edges + \#reversed edges + \#missed edges


## Heatmaps, $\mathrm{n}=1000$

## ER Graphs:



SF Graphs:


## FDR, SHD, n = 1000



Method $\rightarrow$ FGS $\rightarrow$ NOTEARS $\rightarrow$ NOTEARS-L1

## Comparison to global optimum

| $n$ | $\lambda$ | Graph | $F(W)$ | $F\left(W_{\mathrm{G}}\right)$ | $F(\widehat{W})$ |
| ---: | ---: | ---: | ---: | ---: | ---: |
| 20 | 0 | ER2 | 5.11 | 3.85 | 5.36 |
| 20 | 0.5 | ER2 | 16.04 | 12.81 | 13.49 |
| 1000 | 0 | ER2 | 4.99 | 4.97 | 5.02 |
| 1000 | 0.5 | ER2 | 15.93 | 13.32 | 14.03 |
| 20 | 0 | SF4 | 4.99 | 3.77 | 4.70 |
| 20 | 0.5 | SF4 | 23.33 | 16.19 | 17.31 |
| 1000 | 0 | SF4 | 4.96 | 4.94 | 5.05 |
| 1000 | 0.5 | SF4 | 23.29 | 17.56 | 19.70 |


| $\Delta\left(W_{\mathrm{G}}, \widehat{W}\right)$ | $\left\\|\widehat{W}-W_{\mathrm{G}}\right\\|$ | $\left\\|W-W_{\mathrm{G}}\right\\|$ |
| ---: | ---: | ---: |
| -1.52 | 0.07 | 3.38 |
| -0.68 | 0.12 | 3.15 |
| -0.05 | 0.02 | 0.40 |
| -0.71 | 0.12 | 2.95 |
| -0.93 | 0.08 | 3.31 |
| -1.12 | 0.15 | 5.08 |
| -0.11 | 0.04 | 0.29 |
| -2.13 | 0.13 | 4.34 |

- W: ground truth
- W_G: global optimum of NOTEARS M-estimator
- W_hat: Aug. Lagrangian (near) limit point of NOTEARS M-estimator


## Sensitivity to Initialization

Initialization: $W_{\text {init }} \sim \operatorname{Sphere}(r)$ uniformly. What happens if we vary the radius $r \in[0,20]$ ?

(Surprisingly) robust to initialization.

- Erdos-Renyi graph, Linear Gaussian SEM, d=20, num edge =20, $\mathrm{n}=$ \{population, 200\}, L1 regularization parameter $=0.1$
- Each line plot is a random initialization with different $r$.


## Real data example: Cytometry Data

Raw measurement data from Sachs et al. (2005).
Expression levels of proteins and phospholipids in human immune system cells ( $\mathrm{n}=7466 \mathrm{~d}=11$, 20 edges).


## Summary

- We reduce learning DAGs to continuous optimization via a novel M-estimator
- in contrast to conditional independence test based, and local search based methods
- Bridges gap between Markov and Bayesian networks with respect to scalable estimation
- Ongoing Work:
- Analyze landscape of non-convex DAG regularization function: conditions under which it is feasible to get to global optimum
- Approximate fast solvers for matrix exponential

