(Un)expected Encounters of Algebraic Complexity Theory in Machine Learning

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Probabilistic Circuits

Identification in Structural Causal Models





I would like to discuss lower bounds for syntactically multilinear circuits... I would like to discuss lower bounds for syntactically multilinear circuits...

I know everthing about it...

Probabilistic Modelling

Setup:

• Random variables X_1, X_2, \ldots with a joint probability distribution p.

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• State space $\operatorname{val}(X_1) \times \operatorname{val}(X_2) \times \ldots$

Evidence query:

- Given $\xi_i \in \operatorname{val}(X_i)$
- compute $\Pr[X_1 = \xi_1, X_2 = \xi_2, ...]$

Efficiently possible if we can evaluate p efficiently.

Marginal queries

Example:

$$\bullet X_{C} = \begin{cases} 1 & \text{if the train station in city } C \text{ is closed} \\ 0 & \text{otherwise} \end{cases}$$

• joint probability distribution on $Date \times X_C$ _{cities C}

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• What is
$$\Pr[\text{Date} = \text{Mar } 31, X_{\text{Bochum}} = 1]$$
?

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- ▶ We need to "marginalize out" all other cities → exponential sum

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- What is $\Pr[Date = Mar \ 31, X_{Bochum} = 1]$?
- ▶ We need to "marginalize out" all other cities → exponential sum
- Hardness of marginal queries supported by practical evidence

Probabilistic Circuits

Probabilistic circuits (PC)

- compute probablity mass functions
- input nodes are "basic" probability distributions
 - Gaussian distributions (we do not get polynomials)
 - Categorial distributions: $\Pr[X_i = 1]x_i + (1 \Pr[X_i = 1]\bar{x}_i.$
- product nodes
- sum nodes computing weighted sums (with nonnegative weights)

Evidence queries are possible in polynomial times in the circuit size. What about marginal queries?

Decomposable circuits

Definition

A PC is decomposable if the scopes of the children of each product gate are disjoint.

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Theorem

If a distribution is given by a decomposable PC, then we can efficiently perform marginalization.

Probabilistic Generating Circuits

- Given categorical variables X_1, \ldots, X_n with image $\{0, \ldots, d-1\}$
- and joint distribution $p(a_1, \ldots, a_n) = \Pr[X_1 = a_1, \ldots, X_n = a_n]$,
- the *probability generating function* is a formal polynomial in formal variables z_1, \ldots, z_n defined by

$$G(z) = \sum_{j_1=0}^{d-1} \dots \sum_{j_n=0}^{d-1} p(j_1, \dots, j_n) z_1^{j_1} \cdots z_n^{j_n}.$$
 (1)

Definition (Zhang et al. 2021)

A *probabilistic generating circuit (PGC)* for a probability distribution p is an arithmetic circuit that computes G.

- PCs compute probability mass functions
- PGCs store probability distributions as formal objects

Probabilistic Generating Circuits

Theorem (Zhang et al. 2021)

PGCs support efficient marginalization for binary random variables

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PGCs subsume

- PCs
- DPPs (using the algorithm of Mahajan & Vinay).

Determinantal Point Processes (DPP)

Definition

A probability distribution p over n binary random variables X_1, \ldots, X_n is an L-ensemble if there exists a (symmetric) positive semidefinite matrix L such that for all $x \in \{0, 1\}^n$ is distributed according to $\det L_x$, where L_x is the principal minor with the ith rows and columns chosen if $x_i = 1$.

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- L is called the kernel
- \blacktriangleright If L is psd, then all $\det L_x$ are nonnegative
- Normalizing constant is det(L + I).
- Allows for efficient marginalization

Our results

Theorem (Agarwal & B., 2024)

Marginalization is #P-hard for PGC if the categories have size ≥ 3 .

Theorem (Agarwal & B., 2024)

For every PGC over binary variables, there is an equivalent nonmonotone PC (that computes the probability mass function.)

$$g(x_1,\overline{x_1},...,x_n,\overline{x_n}) = f(\frac{x_1}{\overline{x_1}},\frac{x_2}{\overline{x_2}},...,\frac{x_n}{\overline{x_n}}) \cdot \prod_{i=1}^n \overline{x_i}.$$

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Our Results (2)

It is essential how we store the basic distributions:

•
$$\sum_{i=0}^{d-1} \alpha_i x^i$$

• $\sum_{i=0}^{d-1} \alpha_i z_i$

Theorem (Agarwal & B., 2024)

Let C be a nonmonotone PC of size s computing a probability distribution over categorical random variables X_1, \ldots, X_n such that the polynomial P computed by C is set-multilinear with respect to the partition $\{z_{i,0}, \ldots, z_{i,d-1}\}, 1 \leq i \leq n$. Let $A_1, \ldots, A_n \subseteq \{0, \ldots, d-1\}$. Then we can compute $\Pr[X_1 \in A_1, \ldots, X_n \in A_n]$ in time O(s).

ML — ACT dictionary

probabilistic circuit decomposable sum product network subcircuit tree circuit polynomial

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monotone arithmetic circuit syntactically multilinear monotone layered circuit parse tree sum of parse trees representation

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Conclusion: We are not as smart as we think

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Interesting Questions

- Are there more expressive models that are still tractable?
- Are the "generalizations" of DPPs?
- Are there models that subsume DPPs and PCs and compute a nonnegative function *by design*? Formally: Is there a large circuit class such that given C, we can decide in polynomial time whether C is nonnegative on the part of the domain of interest?

Probabilistic Circuits

Identification in Structural Causal Models



Polynomial Identity Testing

We all love PIT!

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Polynomial Identity Testing

We all love PIT!

We investigate algorithms for the problem of Polynomial Identity testing (PIT). Given a polynomial in some implicit representation, it asks whether the polynomial is identically zero or not. It is a fundamental problem in algorithms and complexity theory. It has found applications in algorithm design, for example in algorithms for perfect matching in graphs [17, 36, 39], for primality testing [2, 3, 4], for equivalence testing of read once branching programs [13], and for multi-set equality testing [14], and also in complexity theory, for example, in establishing some major results related to interactive proofs and probabilistically-checkable proofs [38, 9, 8, 7, 45]. In fact, it has also been discovered that an algorithm for polynomial identity testing is intimately connected with complexity theoretic lower bounds [31, 1].

Causation versus Correlation?

Does smoking cause lung cancer?

- Random variable X₁ = number of cigarettes smoked
- $X_1 = f(\varepsilon_1)$ is a function of an unknown random variable ε_1 .
- ▶ Random variable X₂ = binary random variable whether one develops lung cancer.

- $X_2 = g(X_1, \varepsilon_1)$ is a function of X_1 and an unknowm random variable ε_1 .
- Observed correlation between X₁ and X₂.
- Does X_2 cause X_3 or are ϵ_1 and ϵ_2 are correlated.

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Identification in Structural Causal Models

- random variables V_0, V_1, \ldots, V_n
- $\blacktriangleright \ V_i = \sum_j \lambda_{j,i} V_j + \varepsilon_i$ with some additional error term ε_i
- error terms are normally distributed with zero mean
- covariances between the terms given by $\Omega = (\omega_{i,j})$
- models are *recursive*, i.e., for all j > i, we have $\lambda_{j,i} = 0$.



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Identification

Given covariances between the V_i , can we recover the $\lambda_{i,j}$ and $\omega_{i,j}$?

In the example:

•
$$\sigma_{0,1} = \lambda_{0,1}$$
 and $\sigma_{0,2} = \lambda_{0,1}\lambda_{1,2}$. Thus $\lambda_{1,2} = \frac{\lambda_{0,1}\lambda_{1,2}}{\lambda_{0,1}} = \frac{\sigma_{0,2}}{\sigma_{0,1}}$

Generic Identification

- Given a mixed graph G = (V, D, B), (G, D) is acyclic.
- Relation of observed covariances and parameters is given by

$$\boldsymbol{\Sigma} = (\boldsymbol{I} - \boldsymbol{\Lambda})^{-1}\boldsymbol{\Omega}(\boldsymbol{I} - \boldsymbol{\Lambda})^{-T}$$

Definition

G is identifable, if for all Σ (in the image), the parameters Λ (and $\Omega)$ are uniquely determined.

Solved, but somewhat boring.

Definition

G is generically identifiable, if for almost all Σ (in the image), the parameters Λ (and Ω) are uniquely determined.

Methods for Identification

Complexity of the problem is widely open



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- IV = Instrumental variable
- HTC = Halftrek criterion (Foygel, Draisma, Drton 2012)
- ▶ Gröbner basis: Sullivant, Garcia-Puente, Spielvogel 2010
- ETR = Existential theory of the reals (Dörfler et al. 2025)

Hardness of identification

Numerical identification:

• Given a structural causal model and a feasible matrix of observed covariances (Σ).

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• Decide whether there is one ore more solutions (Λ and Ω).

Theorem (Dörfler et al. 2024)

Numerical identification is hard for $\exists \mathbb{R}$.

- ▶ $\exists \mathbb{R} =$ existential theory of the reals.
- still hard if we plant an easy solution.

Tree-shaped Structural Causal Models

Definition (Tree-shaped SCM)

If the directed edges form a directed tree with root 0, then the SCM is called tree-shaped.



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Key feature: Each node has at most one incoming directed edge.

Theorem (Van der Zander, AISTATS 2022)

Generic identification in tree-shaped SCMs is in PSPACE.*

Our result

Theorem (Gupta & B, 2024)

There is an randomized polynomial algorithm that given a tree-shaped SCM M, determines for each parameter $\lambda_{i,j}$ whether it is

- generically identifiable or
- generically 2-identifiable or
- generically unidentifiable.

In the first two cases, it provides corresponding expressions.

Main features:

- polynomial running time
- completeness for tree-shaped SCMs
- 100% PIT

Identification in Tree-shaped SCMs

Theorem (Van der Zander et al., AISTATS 2022)

 $\lambda_{x,y}$ is generically identifiable (k-identifiable) iff the system

$$\begin{split} \lambda_{p,i}\lambda_{q,j}\sigma_{p,q} - \lambda_{p,i}\sigma_{p,j} - \lambda_{q,j}\sigma_{i,q} + \sigma_{i,j} &= 0 \quad \text{for each missing } i \leftrightarrow j \\ \lambda_{p,i}\sigma_{0,p} - \sigma_{0,i} &= 0 \quad \text{for each missing } 0 \leftrightarrow i \end{split}$$

has a unique solution for $\lambda_{x,y}$ (k solutions, respectively).



Möbius transforms

- undirected graph G
- nodes are variables
- edges are labeled with a bilinear equation in the incident variables:

$$axy - bx + cy - d = 0$$

▶ To eliminate y, write

$$y = \frac{bx + d}{ax + c}$$

- This is a Möbius transform. Inverse is given by the adjoint matrix.
- We assume that all $\begin{pmatrix} b & d \\ a & c \end{pmatrix}$ have rank two.
- In the application, rank one edges can be dealt separately. (Rank testing = PIT)

Solving systems of bilinear equations

Let

$$axy - bx + cy - d = 0,$$

$$Ayz - By + Cz - D = 0$$

be two such bilinear equations, sharing the indeterminate y.

To eliminate y, write

$$y = \frac{bx + d}{ax + c}, \quad z = \frac{By + D}{Ay + C}$$

and obtain

$$z = \frac{(Bb + Da)x + (Bd + Dc)}{(Ab + Ca)x + (Ad + Cc)}.$$

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- New coefficients: $\begin{pmatrix} B & D \\ A & C \end{pmatrix} \begin{pmatrix} b & d \\ a & c \end{pmatrix} = \begin{pmatrix} Bb+Da & Bd+Dc \\ Ab+Ca & Ad+Cc \end{pmatrix}$.
- Composition of Möbius transforms



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- Two equations for x of the form $x = \frac{bx+d}{ax+c}$, but they could be trivial.
- If one is nontrivial, solution can be propagated to all other variables.

When is an equation trivial?

Lemma

Let x_1, \ldots, x_t, x_1 be a simple directed cycle and $\begin{pmatrix} b & d \\ a & c \end{pmatrix}$ be the corresponding product.

- 1. If $a \neq 0$, then x_1 has at most two solutions (depending on the discriminant).
- 2. If a = 0 but $c b \neq 0$, then x_1 has exactly one solution.
- 3. If a = c b = 0 but $d \neq 0$, then there is no solution.
- 4. If a = c b = d = 0, then x_1 has infinitely many solutions.
- 1. Solutions given by $\frac{-(c-b)\pm\sqrt{\Delta}}{2a}$. Zero solutions cannot happen here.
- 2. Solutions given by $\frac{d}{c-b}$ (rationally identifiable).
- 3. Cannot happen in our application.
- \longrightarrow 100% PIT

How to find such an identifying cycle?

Van der Zander et al. enumerate all cycles (PSPACE).

Definition

We call a closed walk identifying if $\begin{pmatrix} b & d \\ a & c \end{pmatrix}$ is not a multiple of the identity matrix.

- Let W be the weighted adjacency matrix with 2×2 -matrices as entries.
- Diagonal elements of W^t are the sums of all matrices of closed walks of length t.
- Problem 1: We need simple walks.
- Problem 2: Cancellations

Lemma

If there is an identifying walk of length t, then there is an identifying simple walk of length \leqslant t.

Finger printing



- weight of $(v_i^{(k)}, v_j^{(k+1)})$ is $x_{i,j}^{(k+1)} w(v_i, v_j)$, $w(v_i, v_j)$ weight of original edge.
- ▶ No cancellations. Use PIT to check for an identifying cycle.
- Use self-reduction to find an identifying cycle.

Overall algorithm

Input: A tree-shaped mixed graph M = (V, D, B)

Output: For each $\lambda_{p,i}$, we output whether it is generically identifiable, 2-identifiable, or unidentifiable. In the first two cases, we output corresponding FASTPs.

- 1: Find all rank-1 edges in the missing edge graph (V, \bar{B}) .
- 2: For each missing rank-1 edge $i \leftrightarrow j$, check which of the parameters $\lambda_{p,i}$ or $\lambda_{q,j}$ we can identify. Mark the node i or j, respectively.
- 3: Remove all rank-1 edges from the missing edge graph. Call the resulting graph H. Let C_1, \ldots, C_t be the connected components of H.
- 4: for each connected component $C_i \mbox{ do}$
- 5: **if** C_i contains a marked node **then**
- 6: Propagate the result to all unidentified nodes in C_i .
- 7: **else**
- 8: Find an identifying cycle in C_i (using PIT).
- 9: If no such cycle is found, report that all nodes of C_i are unidentifiable.
- 10: If the cycle produces one solution, then progagate it to all the nodes of C_i .
- 11: If the cycle produces two solutions, then propagate it to all the nodes of C_i .
- 12: Plug the solutions into the equations of C_i and use PIT to check whether all equations are satisfied.

Implementation

We benchmarked our code on 856 SCMs with 8 nodes (from van der Zander et al.)



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computational complexity

- Since 2025, I am the editor-in-chief of computational complexity.
- Founded by Joachim von zur Gathen in the spirit of Schönhage and Strassen.
- Transformative journal.
- Editorial board contains 5 Turing award winners, 2 Abacus prize winners, 1 Abel prize winner.
- Quantum is covered, too: Thomas Vidick and Francois Le Gall.
- If mathematicians (and physicists) are to take us seriously, we need to submit to journals, too.

Please consider submitting very good papers of yours!