Partial Homoscedasticity in Causal Discovery with Linear Models

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• Linear SEM: random vector X solves

$$X = \Lambda^T X + \epsilon,$$
 $Var[\epsilon] = diag(\omega).$

- Represented by a directed ayclic graph (DAG): G = (V, E)
- The DAG / SEM has a covariance parameterization:

$$\begin{split} \phi_{\mathcal{G}} : \mathbb{R}^{\mathcal{E}} \times \mathbb{R}^{\mathcal{V}} &\mapsto PD, \\ (\Lambda, \boldsymbol{\omega}) &\mapsto (I - \Lambda)^{-\mathcal{T}} \mathsf{diag}(\boldsymbol{\omega})(I - \Lambda)^{-1}, \end{split}$$

with

$$\mathbb{R}^{E} = \left\{ \Lambda \in \mathbb{R}^{V \times V} : \lambda_{ij} = 0 \text{ if } i \to j \notin E \right\}.$$

Example 1

$$X_{1} = \varepsilon_{1}$$

$$X_{2} = \varepsilon_{2}$$

$$X_{3} = \lambda_{13}X_{1} + \lambda_{23}X_{2} + \varepsilon_{3}$$

$$X_{4} = \lambda_{24}X_{2} + \lambda_{34}X_{3} + \varepsilon_{4}$$

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$$M = \begin{pmatrix} 0 & 0 & \lambda_{13} & 0 \\ 0 & 0 & \lambda_{23} & \lambda_{24} \\ 0 & 0 & 0 & \lambda_{34} \\ 0 & 0 & 0 & 0 \end{pmatrix}, \qquad \omega = (\omega_{1}, \omega_{2}, \omega_{3}, \omega_{4})^{T}.$$

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Structural Identifiability

- Classic case: Markov equivalence classes of DAGs
- Homoscedastic errors: if all error variances are equal, the DAG *G* is uniquely identifiable

[Chen, Drton, and Wang 2019; Peters and Bühlmann 2014]

- Homoscedastic error, general directed graphs?
- Setup in between: partial homoscedasticity (groupwise equal error variances)

Our contributions

- Derive an implicit description of linear Gaussian SEM under partial homoscedasticity
- Characterize the DAGs that define the same partially homoscedastic linear Gaussian SEM
- Give an algorithm for constructing equivalence class representation

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 A partition of nodes w.r.t. different error variances ω₁,..., ω_K: Π = {π₁,..., π_K}. The k'th (homoscedastic errors) block of nodes is the set π_k = {i₁, i₂,..., i_m} such that

$$\omega_{i_1i_1}=\omega_{i_2i_2}=\cdots=\omega_{i_mi_m}=\omega_k.$$

- For i, j in the same block: $i \sim_{\Pi} j$.
- The groupwise homoscedastic linear Gaussian model given by G = (V, E) and Π :

$$M_{G,\Pi} = \left\{ \Sigma : \Sigma = \phi_G(\Lambda, \boldsymbol{\omega}), \ \Lambda \in \mathbb{R}^E \text{ and } \boldsymbol{\omega} \in \mathbb{R}^V, \omega_{ii} = \omega_{jj} \text{ if } i \sim_{\Pi} j \right\}$$

Theorem 1 (Drton 2018)

The error variance ω_{ii} can be computed from the true covariance matrix

$$\omega_{ii} = \sigma_{ii} - \Sigma_{i,\mathcal{A}} (\Sigma_{\mathcal{A},\mathcal{A}})^{-1} \Sigma_{\mathcal{A},i},$$

for any A such that $pa(i) \subseteq A \subseteq V \setminus de(i)$.

• As a corollary, if *i* and *j* are in the same block then

$$\sigma_{ii} - \Sigma_{i,\mathcal{A}_i} (\Sigma_{\mathcal{A}_i,\mathcal{A}_i})^{-1} \Sigma_{\mathcal{A}_i,i} = \sigma_{jj} - \Sigma_{j,\mathcal{A}_j} (\Sigma_{\mathcal{A}_j,\mathcal{A}_j})^{-1} \Sigma_{\mathcal{A}_j,j}$$

holds for all A_i and A_j such that $pa(i) \subseteq A_i \subseteq V \setminus de(i)$ and $pa(j) \subseteq A_j \subseteq V \setminus de(j)$.

Theorem 2

If i and j are in the same block of partition, then the constraint equation

$$\sigma_{ii} - \Sigma_{i,\mathcal{A}_i} (\Sigma_{\mathcal{A}_i,\mathcal{A}_i})^{-1} \Sigma_{\mathcal{A}_i,i} = \sigma_{jj} - \Sigma_{j,\mathcal{A}_j} (\Sigma_{\mathcal{A}_j,\mathcal{A}_j})^{-1} \Sigma_{\mathcal{A}_j,j}$$

holds for all matrices $\Sigma \in M_{G,\Pi}$ if and only if

$$\operatorname{pa}(i) \subseteq A_i \subseteq V \setminus \operatorname{de}(i) \text{ and } \operatorname{pa}(j) \subseteq A_j \subseteq V \setminus \operatorname{de}(j).$$

The partially homoscedastic linear Gaussian model $M_{G,\Pi}$ is uniquely determined by

- $\bullet\,$ Equal variance constraints induced by $\Pi\,$
- Conditional independence constraints induced by G

Partial Homoscedasticity

Example: $\Pi = \{\{1, 2\}, \{3\}\}$



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$$\mathcal{M}_1 = \{ \Sigma \mid \underbrace{\sigma_{11}\sigma_{33} = \sigma_{22}\sigma_{33} - \sigma_{23}^2}_{\operatorname{Var}[X_2|X_3] = \operatorname{Var}[X_1]}, \underbrace{\sigma_{13}\sigma_{23} - \sigma_{12}\sigma_{33} = 0}_{X_1 \perp \perp X_2|X_3}, \Sigma \in \mathcal{PD}_3 \}$$

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$$\mathcal{M}_2 = \{ \Sigma \mid \underbrace{\sigma_{22}\sigma_{33} = \sigma_{11}\sigma_{33} - \sigma_{13}^2}_{\operatorname{Var}[X_1|X_3] = \operatorname{Var}[X_2]}, \underbrace{\sigma_{13}\sigma_{23} - \sigma_{12}\sigma_{33} = 0}_{X_1 \perp \perp X_2 \mid X_3}, \Sigma \in \mathcal{PD}_3 \}$$

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Theorem 3

Given a partition Π of nodes, two DAGs G_1, G_2 have the same partially homoscedastic linear Gaussian model ($M_{G_1,\Pi} = M_{G_2,\Pi}$) if and only if the following two conditions hold:

- **(**) G_1 and G_2 have the same skeleton and unshielded colliders.
- ② For each node *i* with *i* ∈ π_k , $|\pi_k| \ge 2$, the parents of *i* in *G*₁ and *G*₂ are the same: pa₁(*i*) = pa₂(*i*).

In this case we say that G_1 and G_2 are (distributionally) equivalent given the partition Π : $G_1 \approx_{\Pi} G_2$.

Two extreme cases:

- $\Pi = \Pi_{\text{finest}} = \{\{i\} : i \in V\}$: classic setup, all different variances
- $\Pi = \Pi_{\text{coarsest}} = \{V\}$: all equal variance, uniquely identifiable

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Definition: CPDAG

A completed partially directed acyclic graph (CPDAG) of a DAG G under partition Π is defined as the union of all equivalent DAGs of G:

 $G^*_{\Pi} := \cup (G' \mid G' \approx_{\Pi} G).$



• Just the classic CPDAG definition, but the construction is a bit different!

The Orientation rules R1-R4 in [Meek 1995], for propagating orientations



Figure: The 4 Orientation rules.

Algorithm Constructing the equivalence class of a DAG, given the partition

Require: A DAG G, the partition Π

- 1: Create an empty graph G'
- 2: Copy the skeleton and all edge orientations with unshielded colliders of G to G'
- 3: Apply rules R1, R2 and R3 on G' until no more edges can be oriented
- 4: for $i \in V$ with $i \in \pi_k$ and $|\pi_k| \ge 2$ do
- 5: Copy the orientation of edges in G having one endpoint at i to G'
- 6: end for
- 7: Apply rules R1 and R2 on G' until no more edges can be oriented
- 8: return $G' = G_{\Pi}^*$

Equivalence Classes



Figure: A DAG and the corresponding CPDAG, under a fixed partition.

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- Extension of equal variance assumption
- New perspective of model identifibility

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- Restrictive conditions
- Cannot be applied to cycles

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