Causal Inference With Contagion and Latent Homophily Under Full Interference

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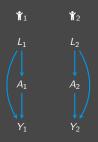
Williams College

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L = confounders; A = therapy sessions; Y = job satisfaction.

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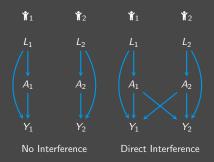
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No Interference

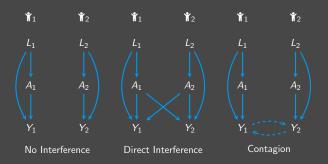
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Social Interactions Create Dependence in Data $^{ m 1}$

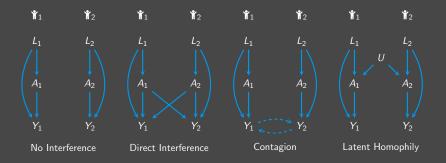


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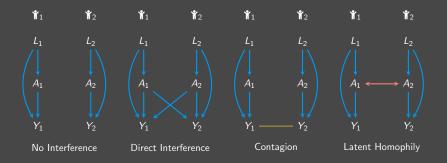


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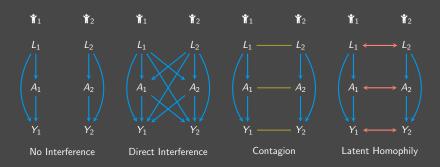


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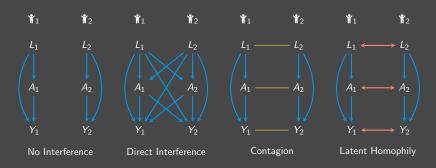
Social Interactions Create Dependence in Data $^{ m 2}$



²Shalizi and Thomas 2011, Ogburn and VanderWeele 2014, Lauritzen and Richardson 2002, Shpitser 2015



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[&]quot;Interference"

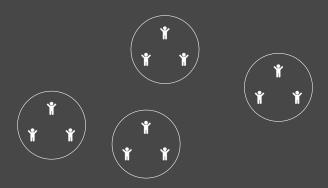
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Partial Interference: assume i.i.d. "chunks" of data. 4

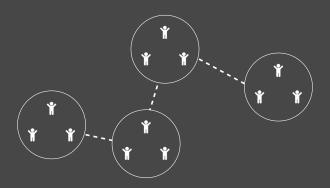
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Partial interference does not hold in general!



This work focuses on **Full Interference** ⁵:

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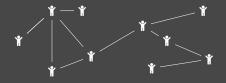
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Usually assume **parameter-sharing**: $p(L_i, L_{\mathsf{nb}(i)}; \theta)$ is shared by everyone in the network.

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Auto-g computation⁶:

► can estimate causal effects under full interference.

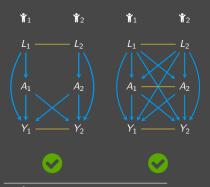
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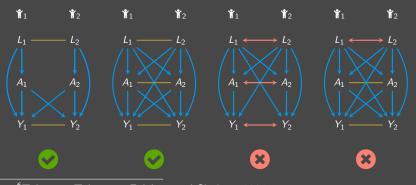
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Causal Inference for Social Network Data 7.

allows for direct interference and latent homophily between individuals.

Open Problem

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Bidirected Edge:

$$Y_1 \longleftrightarrow Y_2 \longleftrightarrow Y_3$$

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How To Get i.i.d. Samples for Independence Tests

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Our Proposed Test

Step 1: find a 5-hop independent set \mathcal{I} from the network.



Our Proposed Test

Step 2: for each person in \mathcal{I} , collect information on their neighbors and their 2nd-order neighbors (i.e., neighbors' neighbors).



Our Proposed Test

Step 3: Is $person_i \perp \!\!\! \perp 2nd$ -order $nb(i) \mid nb(i)$?

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Likelihood ratio test:

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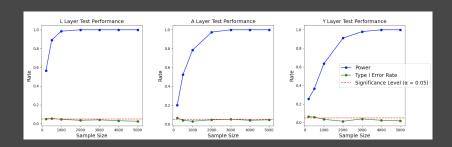
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```
If \perp \!\!\! \perp, conclude contagion (-).
```

If $\not\perp$, conclude latent homophily (\leftrightarrow) .

Evaluating Our Test



Recap

- A causal effect estimation method that allows for three types
 of dependence mechanisms: direct interference (→),
 contagion (−), and latent homophily (↔).
- The assumptions of auto-g relies on prior knowledge & belief.
 We want a test to distinguish between interference due to contagion (–) and latent homophily (↔).

Why do we even need a new method when latent homophily (\leftrightarrow) is present?

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Undirected Edge:

$$L_1$$
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Gibbs factors ⁸: $p(L_1 \mid L_2)$, $p(L_2 \mid L_1, L_3)$, and $p(L_3 \mid L_2)$

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Bidirected Edge:

$$\begin{array}{ccc} \mathsf{Cov}(1,2) & \mathsf{Cov}(2,3) \\ \mathsf{L}_1 & \longleftrightarrow & \mathsf{L}_2 & \longleftrightarrow & \mathsf{L}_3 \end{array}$$

$$p(L_1, L_2, L_3) \sim MVN(\mu, \Sigma)$$

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Target for Causal Effect Estimation

Unit potential outcome expectation⁹ for every $i \in V$:

$$\mathbb{E}[Y_i(a)] = \sum_{l} \mathbb{E}[Y_i \mid A = a, L] \times p(L)$$

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Latent Homophily (\leftrightarrow) :

1. Pre-processing: select triplets from the network s.t. none of these triplets are connected.



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- 2. Assume $p(L) \sim MVN$ & Parameter-sharing.
- 3. Apply Residual Iterative Conditional Fitting (RICF) 10 : $\widehat{\mu}$, $\widehat{\Sigma}$.

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Task 2: Estimate $\mathbb{E}[Y_i \mid A, L]$

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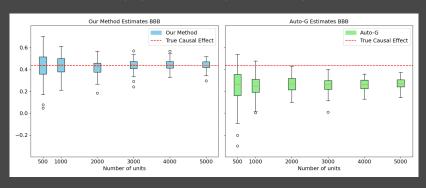
Latent Homophily (\leftrightarrow) :



Directly estimate $\mathbb{E}[Y_i \mid A, L]$ using data from people in an independent set of the network.

Latent homophily (\leftrightarrow) in all three (L, A, and Y) layers.

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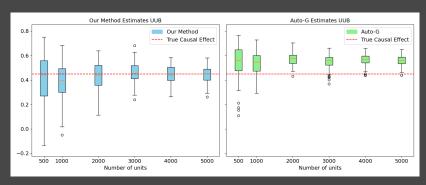


Contagion (-) in the L and A layers.

Latent homophily (\leftrightarrow) in the Y layers.

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Latent homophily (\leftrightarrow) in the Y layers.



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Thanks!

Questions?