

# Causal inference for infectious disease intervention in inter-connected clusters

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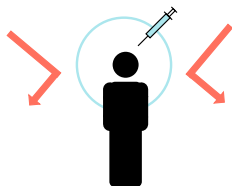
Joint Statistical Meeting 2022

(References and slides are available on my personal website.)

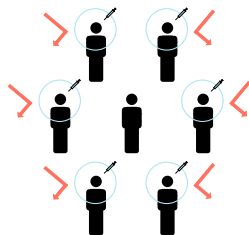
# Infectious disease and vaccination

- Direct protection for the treated individuals:
  - direct effect, vaccine efficacy, susceptibility effect, ...
- Indirect protection for the surrounding individuals:
  - indirect effect, herd immunity, contagion effect, infectiousness effect, ...

(Examples: vaccines for Polio, Influenza, HIV/AIDS, Malaria, and etc).



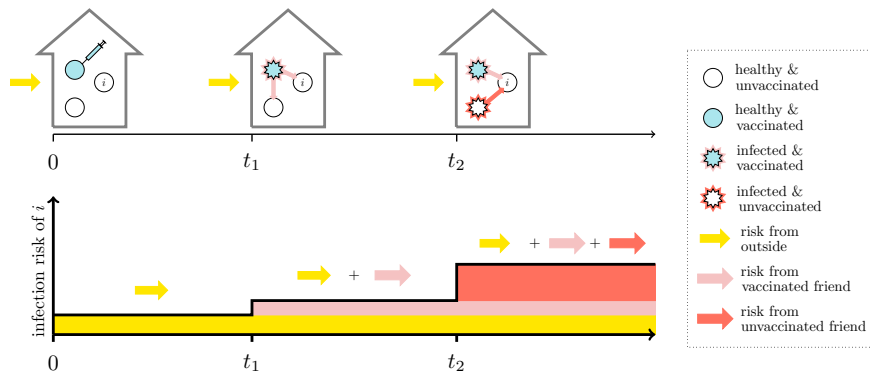
Direct protection



Indirect protection

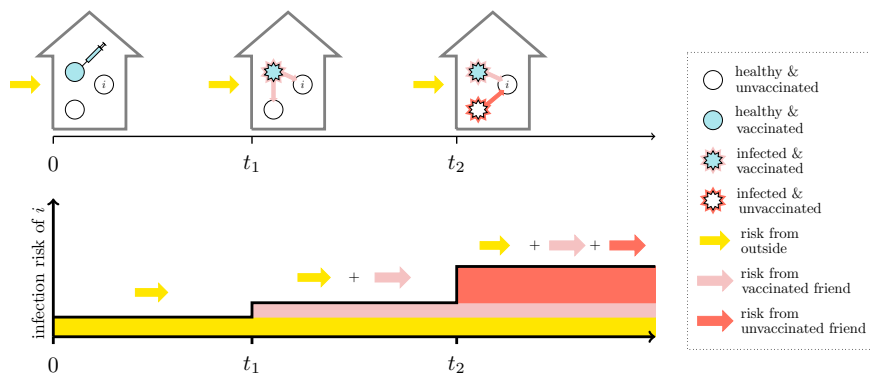
# How epidemiologists understand disease transmission

For a focal individual  $i$ , the risk of infection increases as more neighbors become infectious and depend on neighbors' vaccination status.



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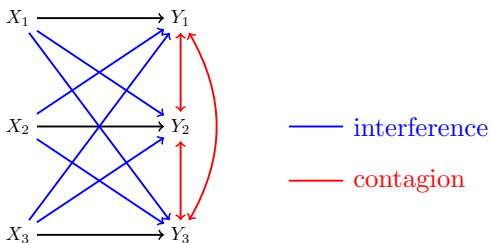


One infection outcome depends on (i) its **own treatment**, (ii) **treatments of others**, and (iii) **infection times of others** (e.g.  $t_1$ ,  $t_2$ ).

# Why is infectious disease difficult to study in causal inference?

## 1 Interdependence of outcomes and treatments across subjects

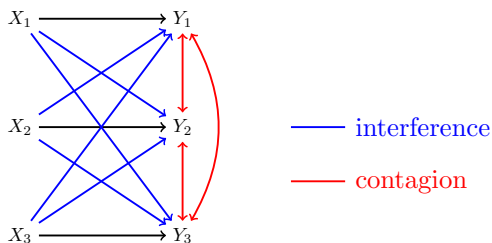
Consider a interconnected three individuals with treatments  $(X_1, X_2, X_3)$  and infection outcomes  $(Y_1, Y_2, Y_3)$ .



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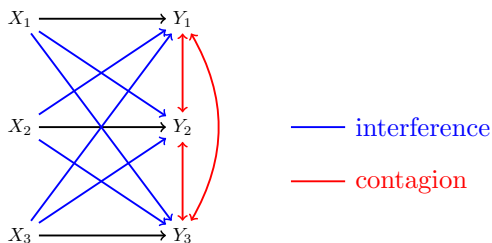


- The infection outcome of one individual depends on others' treatments
- The infection outcome of one individual depends on other's outcomes, since it is transmissible.

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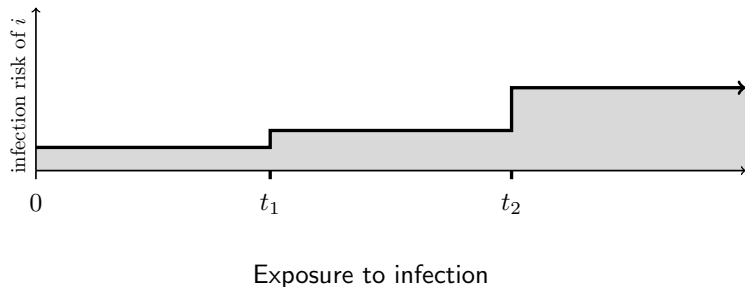


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**Bidirectional arrows causes problems in causal identification**

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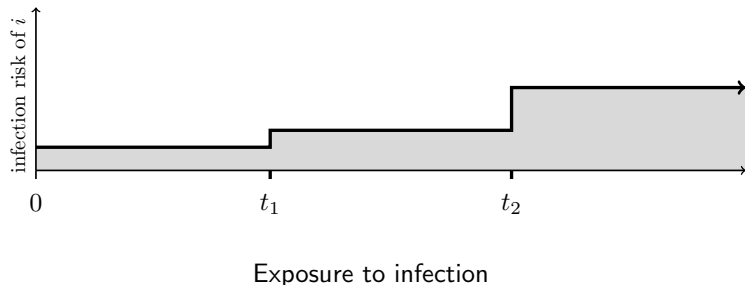
- 1 Interdependence of outcomes and treatments across subjects
- 2 Stochastic processes of “exposure to infection”
  - ▶ “Exposure to infection” is determined by stochastic infection outcomes of others, whose distribution depends on their treatments.





# Why is infectious disease difficult to study in causal inference?

- 1 Interdependence of outcomes and treatments across subjects
- 2 Stochastic processes of “exposure to infection”



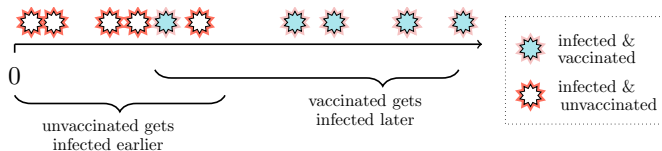
For example, earlier infections of neighbors (smaller  $t_1$  and  $t_2$ ) or higher infectiousness due to unvaccination (bigger jumps) increases overall “exposure to infection” and consequently infection risk.

# Why is infectious disease difficult to study in causal inference?

- 1 Interdependence of outcomes and treatments across subjects
- 2 Stochastic processes of “exposure to infection”
- 3 Bias due to differential “exposure to infection”

Can we directly compare treated and untreated individuals using randomization?

$$E[Y_i|X_i = 1] - E[Y_i|X_i = 0]$$



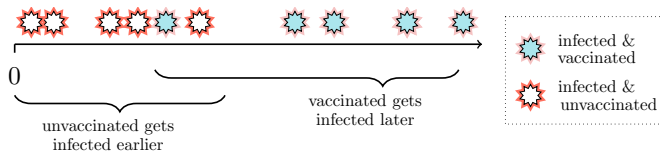
For example, vaccinated individuals endure higher exposure to infection, which is not a fair comparison. → Effect is under-estimated!

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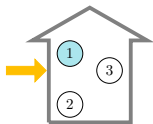


For example, vaccinated individuals endure higher exposure to infection, which is not a fair comparison. → Effect is under-estimated!

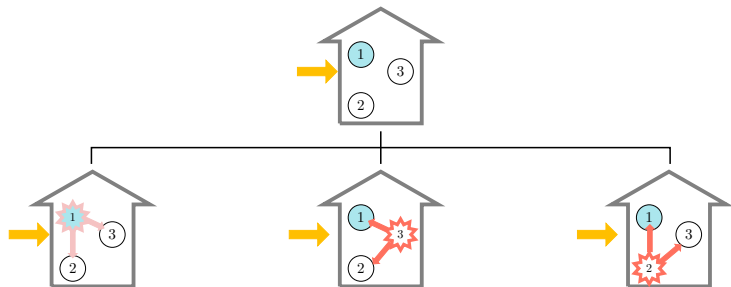
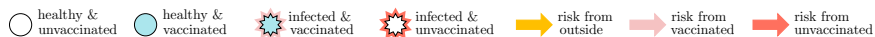
**NO! It may be biased due to differential “exposure to infection”. [1,2,3].**

So how to solve the causal identification problem  
for infectious disease outcomes?

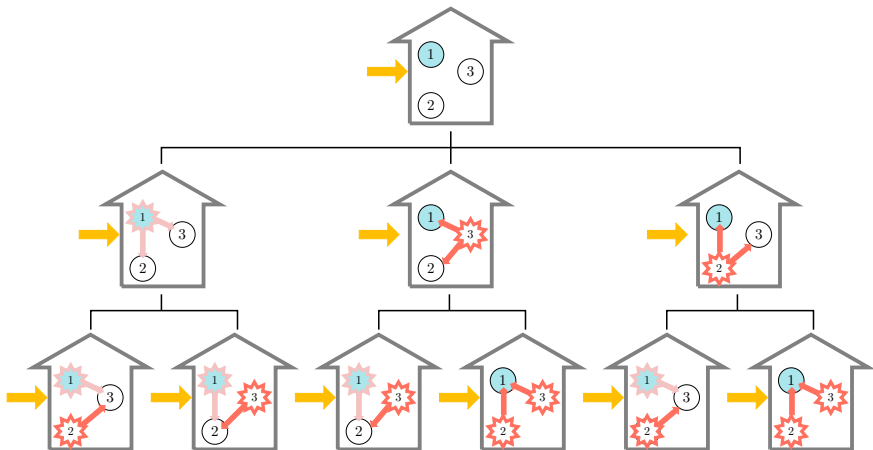
# Decompose infection process regarding different orders



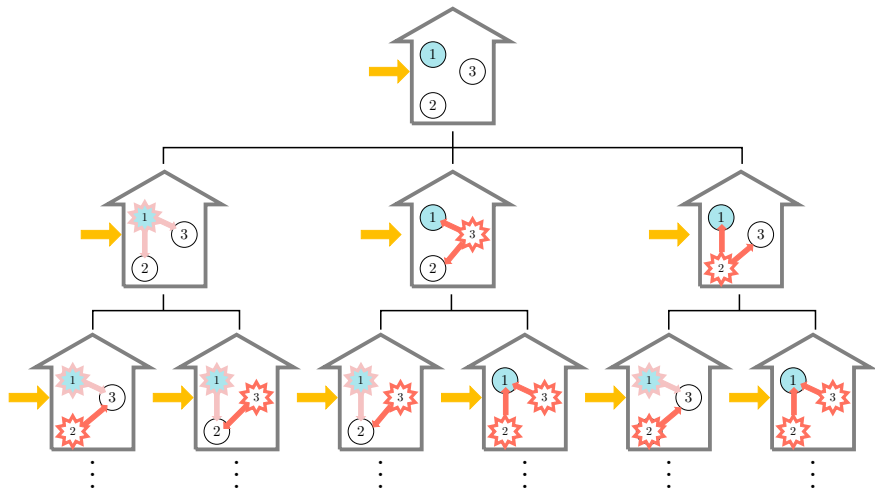
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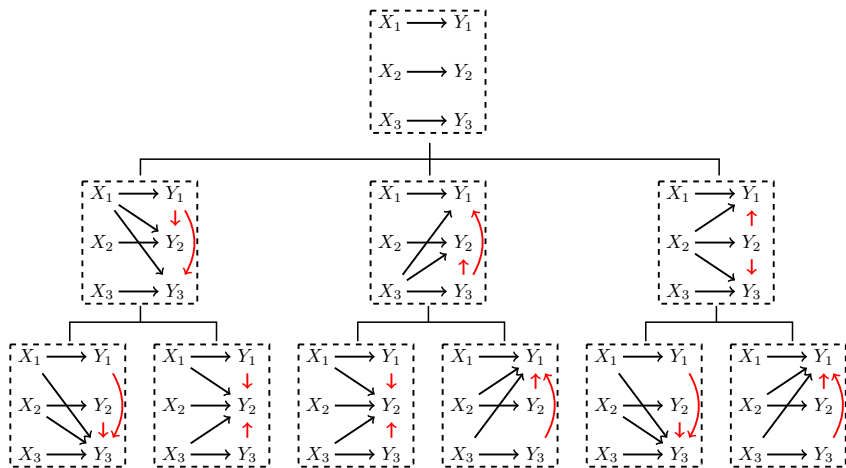
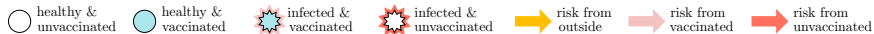


# Decompose infection process regarding different orders





# Decompose infection process regarding different orders



# Notation

- Subject  $i$ 's treatment:  $X_i$
- Others' treatments:  $\mathbf{X}_{(i)} = (X_1, \dots, X_{i-1}, X_{i+1}, \dots, X_n)$
- Others' infection history:  $\mathcal{H}_{(i)}(\mathbf{x}_{(i)}) = \{T_j(\mathbf{x}_{(i)}) : j \neq i\}$

## Potential outcome

Identify  $Y_i(t; x_i, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}(\mathbf{x}_{(i)}))$  as the counterfactual infection outcome at time  $t$  under joint treatment  $(x_i, \mathbf{x}_{(i)})$  and infection times of others  $\mathcal{H}_{(i)}(\mathbf{x}_{(i)})$  under treatments  $\mathbf{x}_{(i)}$ ,

- (i) own treatment:  $X_i = x_i$
- (ii) others' treatments:  $\mathbf{X}_{(i)} = \mathbf{x}_{(i)}$
- (iii) others' infection times:  $\mathcal{H}_{(i)}(\mathbf{x}_{(i)})$

# Identification of exposure-marginalized potential outcomes

## Causal identification

Under conventional assumptions in causal inference, the potential outcome

$\mathbb{E}[Y_i(t; x_i, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}^*(\mathbf{x}'_{(i)})) | \mathbf{L} = \mathbf{l}]$  can be identified as

$$\mathbb{E}[Y_i(t; x_i, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}^*(\mathbf{x}'_{(i)})) | \mathbf{L} = \mathbf{l}] = \int \mathbb{E}[Y_i(t; x_i, \mathbf{x}_{(i)}, \mathbf{h}_{(i)}) | \mathbf{L} = \mathbf{l}] dG_{(i)}^*(\mathbf{h}_{(i)} | \mathbf{x}'_{(i)}, \mathbf{l}_{(i)})$$

where

$$\mathbb{E}[Y_i(t; \mathbf{h}_{(i)}, \mathbf{x}) | \mathbf{L} = \mathbf{l}] = \sum_{j=0}^{n-1} \left[ F_{I_i}^{j+1}(\min\{t, t_{(i)}^{j+1}\} - t_{(i)}^j | \mathbf{x}, \mathbf{h}_{(i)}, \mathbf{l}) \prod_{k=0}^{j-1} (1 - F_{I_i}^k(t_{(i)}^{k+1} - t_{(i)}^k | \mathbf{x}, \mathbf{h}_{(i)}, \mathbf{l})) \right]$$

$$dG_{(i)}^*(\mathbf{h}_{(i)} | \mathbf{x}_{(i)}, \mathbf{l}) = \prod_{j=1}^{n-1} \left[ f_{\varphi_i^j}^{j-1}(t_{(i)}^j - t_{(i)}^{j-1} | \mathbf{x}, \mathbf{h}_{(\varphi_i^j)}, \mathbf{l}) \prod_{k=j+1}^{n-1} S_{\varphi_i^k}^{j-1}(t_{(i)}^j - t_{(i)}^{j-1} | \mathbf{x}, \mathbf{h}_{(\varphi_i^k)}, \mathbf{l}) \right]$$

$$F_{I_i}^k(s | \mathbf{x}, \mathbf{h}_{(i)}, \mathbf{l}) = 1 - \exp \left[ - \int_{t_{(i)}^k}^{t_{(i)}^k + s} \frac{f_i^k(u | \mathbf{x}, \mathbf{h}_{(i)}, \mathbf{l})}{S_i^k(u | \mathbf{x}, \mathbf{h}_{(i)}, \mathbf{l})} du \right] \text{ for } k = 0, \dots, n-1$$

# Exposure-marginalized (natural) causal estimands

## Exposure-marginalized (natural) causal estimands

- Susceptibility effect

$$SE_i(t, \mathbf{x}_{(i)}) = \mathbb{E}[Y_i(t; \mathbf{1}, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}^*(\mathbf{x}_{(i)})) - Y_i(t; \mathbf{0}, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}^*(\mathbf{x}_{(i)}))]$$

- Infectiousness effect

$$IE_i(t, x_i, \mathbf{x}_{(i)}) = \mathbb{E}[Y_i(t; x_i, \mathbf{1}, \mathcal{H}_{(i)}(\mathbf{x}_{(i)})) - Y_i(t; x_i, \mathbf{0}, \mathcal{H}_{(i)}(\mathbf{x}_{(i)}))]$$

- Contagion effect

$$CE_i(t, x_i, \mathbf{x}_{(i)}, \mathbf{x}'_{(i)}) = \mathbb{E}[Y_i(t; x_i, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}^*(\mathbf{x}_{(i)})) - Y_i(t; x_i, \mathbf{x}_{(i)}, \mathcal{H}_{(i)}^*(\mathbf{x}'_{(i)}))]$$

- Susceptibility effect → shows if the vaccine protects treated individual
- Infectiousness effect → shows if the vaccine decreases transmission ability
- Contagion effect → shows if the disease is contagious

# Traditional causal estimands in cluster studies

- Direct effect:

$$DE(t) = \mathbb{E}[Y_i(t)|X_i = 1] - \mathbb{E}[Y_i(t)|X_i = 0]$$

- Indirect effect:

$$\begin{aligned} IDE(t) &= \sum_{|\mathbf{x}_{(i)}|=\frac{n}{2}} \mathbb{E}[Y_i(t)|X_i = 0, \mathbf{X}_{(i)} = \mathbf{x}_{(i)}]p(\mathbf{x}_{(i)}) \\ &\quad - \sum_{|\mathbf{x}_{(i)}|=0} \mathbb{E}[Y_i(t)|X_i = 0, \mathbf{X}_{(i)} = \mathbf{x}_{(i)}]p(\mathbf{x}_{(i)}) \end{aligned}$$

[1] Longini et al. Statistical inference for infectious diseases: risk-specific household and community transmission parameters. *American Journal of Epidemiology*, 128(4):845–859, 1988.

[2] Halloran et al. Direct and indirect effects in vaccine efficacy and effectiveness. *American Journal of Epidemiology*, 133(4):323–331, 1991.

[3] Halloran et al. Exposure efficacy and change in contact rates in evaluating prophylactic HIV vaccines in the field. *Statistics in Medicine*, 13(4):357–377, 1994.

## Simulation: true values of causal estimands

| Cluster | Treatment | Causal estimands |        |        |         |          |
|---------|-----------|------------------|--------|--------|---------|----------|
|         |           | $CE$             | $SE$   | $IE$   | $DE(t)$ | $IDE(t)$ |
| 2       | Obs.      | 0.004            | -0.013 | -0.037 | -0.012  | -        |
|         | Bernoulli | 0.004            | -0.013 | -0.037 | -0.012  | -0.037   |
|         | Block     | -                | -      | -      | 0.024   | -        |
|         | Cluster   | -                | -      | -      | -0.049  | -        |
| 4       | Obs.      | 0.026            | -0.013 | -0.084 | -0.011  | -        |
|         | Bernoulli | 0.026            | -0.013 | -0.084 | -0.011  | -0.063   |
|         | Block     | -                | -      | -      | 0.018   | -        |
|         | Cluster.  | -                | -      | -      | -0.098  | -        |
| 8       | Obs.      | 0.066            | -0.013 | -0.129 | -0.010  | -        |
|         | Bernoulli | 0.066            | -0.013 | -0.129 | -0.010  | -0.094   |
|         | Block     | -                | -      | -      | 0.010   | -        |
|         | Cluster   | -                | -      | -      | -0.151  | -        |

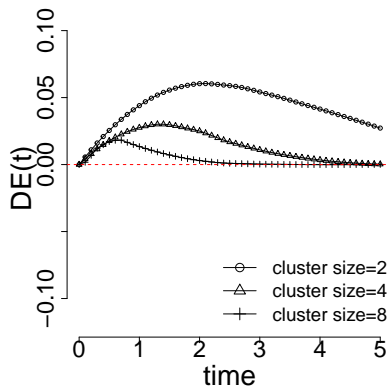
Simulation under  $e^{\beta_1} = 0.9$ ,  $e^{\beta_2} = 0.1$ ,  $\alpha(t) = 0.3$ ,  $\gamma(t) = 3$  and  $e^{\theta_1} = e^{\theta_2} = 0.9$ . Clusters of 2, 4, and 8 are observed at 0.4, 0.3 and 0.2 year.

## Simulation: true values of causal estimands

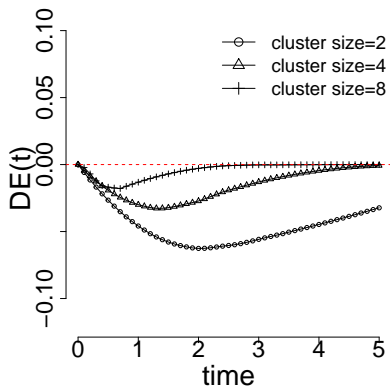
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|         | Cluster   | -                | -       | -       | -0.151  | -        |

Simulation under  $e^{\beta_1} = 0.9$ ,  $e^{\beta_2} = 0.1$ ,  $\alpha(t) = 0.3$ ,  $\gamma(t) = 3$  and  $e^{\theta_1} = e^{\theta_2} = 0.9$ . Clusters of 2, 4, and 8 are observed at 0.4, 0.3 and 0.2 year.

# Simulation: correlated treatment assignment under null direct protection



Negatively correlated



Positively correlated

Simulation under  $e^{\beta_1} = 1$ ,  $e^{\beta_2} = 0.1$ ,  $\alpha(t) = 0.3$ , and  $\gamma(t) = 3$ .



# Estimation

| Treatment | $\hat{CE}$    | $\hat{SE}$      | $\hat{IE}$      | $\hat{DE}(t)$   | $\hat{IDE}(t)$  |
|-----------|---------------|-----------------|-----------------|-----------------|-----------------|
| Obs.      | 0.265         | -0.075          | -0.047          | -0.105          | -               |
|           | (0.189,0.345) | (-0.122,-0.033) | (-0.088,-0.013) | (-0.150,-0.036) | -               |
| Bernoulli | 0.257         | -0.077          | -0.046          | -0.102          | -0.147          |
|           | (0.182,0.335) | (-0.124,-0.035) | (-0.088,-0.013) | (-0.151,-0.052) | (-0.231,-0.060) |
| Block     | -             | -               | -               | -0.065          | -               |
|           | -             | -               | -               | (-0.094,-0.038) | -               |
| Cluster   | -             | -               | -               | -0.448          | -               |
|           | -             | -               | -               | (-0.566,-0.328) | -               |

Estimation with 100 clusters of size 10 at  $t=0.5$ , when  $e^{\beta_1} = 0.5$ ,  $e^{\beta_2} = 0.5$ ,  $\alpha(t) = 0.3$ ,  $\gamma(t) = 3$  and  $e^{\theta_1} = e^{\theta_2} = 0.9$ . True values are  $SE(t = 0.5) = 0.265$ ,  $IE(0.5) = -0.075$ ,  $CE(t = 0.5) = -0.048$  across various treatment assignments, and  $IDE(t) = -0.151$  under Bernoulli randomization. True values of  $DE(0.5)$  are -0.107, -0.107, -0.067, and -0.458, respectively for Obs, Bernoulli, Block, and Cluster randomization.

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| Block     | 0.255<br>(0.179,0.332) | -0.076<br>(-0.122,-0.035) | -0.047<br>(-0.091,-0.012) | -0.065<br>(-0.094,-0.038) | -<br>-                    |
| Cluster   | 0.253<br>(0.150,0.359) | -0.079<br>(-0.146,-0.024) | -0.050<br>(-0.101,-0.011) | -0.448<br>(-0.566,-0.328) | -<br>-                    |

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# Summary

- We articulate the causal structure between individuals' treatments and outcomes of infectious disease in a inter-connected cluster.
- We illustrate the identification strategy for infectious disease potential outcomes non-parametrically
  - ▶ Do not depend on certain study design or randomization strategy
  - ▶ Apply to various transmission dynamics, cluster size and observational time
  - ▶ Incorporate individual- and cluster-level covariates
- A class of novel and biologically meaningful causal estimands for the susceptibility, infectiousness and contagion effect of vaccines are proposed
- We provide comprehensive comparisons to popular estimands in contemporary epidemiology

## Other relevant work and future direction

- We further apply a generalized Cox-type transmission hazard model to facilitate the inference of causal estimands parametrically or semi-parametrically.
- We promote hazard ratio as alternative causal estimands for the susceptibility and infectiousness effect, and compared them to existing estimands for vaccine efficacy.
- Extend current research on causal identification for contagious outcomes to more realistic scenarios, for example, relaxing requirement on accurate infection times, accommodating incomplete knowledge of transmission network, allowing recovering and re-infection of outcomes.

## References

Xiaoxuan Cai, Wen Wei Loh, Forrest W. Crawford. (2021) Identification of Causal intervention effects under contagion. *Journal of Causal Inference*, 9, 9-38. (Winner of best paper award, ASA Section on Statistics in Epidemiology)

Xiaoxuan Cai, Eben Kenah, Forrest W. Crawford. (2021) Causal identification of infectious disease intervention effects in a clustered population. [arXiv:2105.03493](https://arxiv.org/abs/2105.03493)

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<https://xiaoxuan-cai.github.io/>

(References and slides are available on my personal website.)

Thank you!