

# Causal inference for infectious disease intervention under contagion

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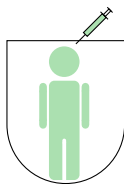
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<https://arxiv.org/abs/1912.04151>

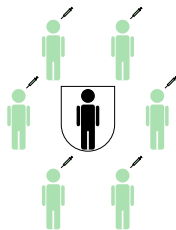
# Infectious disease and vaccination

Distinct mechanisms of infectious disease interventions/vaccinations,

- Direct protection for the treated individuals:
  - direct effect, vaccine efficacy, susceptibility effect...
- Indirect protection for the surrounding individuals:
  - indirect effect, herd immunity...
- Vaccines for Polio, Influenza, HIV/AIDS, Malaria and etc.



Direct protection



Indirect protection

# Why infectious disease is difficult to study?

Research on transmission of infectious disease has some unique features and challenges.

- The infection outcome of one individual also depends on others' treatments, conditional on other individuals being infected.  
– **Interference**
- The outcome of interest (infection) is transmissible, so outcomes are not independent from each other. – **Contagion**
- The infection times of others compose an important factor for the infection outcome – **Exposure to infection**  
*Earlier exposure to infectious individual (higher "exposure to infection") increases the risk of infection.*

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

## Bias due to differential “exposure to infection”

Direct comparisons of treated and untreated individuals may not be valid due to differential “exposure to infection” [1,2,3].

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

- For example, if vaccinated individuals get infected later in general, then later infected and vaccinated subjects face higher exposure to infection, comparing to earlier infected and unvaccinated individuals.  
→ not a fair comparison !

Can randomization solve the problem?

- No. Differential “exposure to infection” due to others’ infection happens after randomization.

[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.

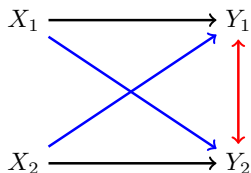
[2] Halloran et al. Direct and indirect effects in vaccine efficacy and effectiveness. *American Journal of Epidemiology*, 133(4):323–331, 1991.

[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.

## Symmetric partnership models

Partnership models have been widely understood as a useful framework to clarify causal relationship in epidemiology, and lay the foundation for more complex settings.

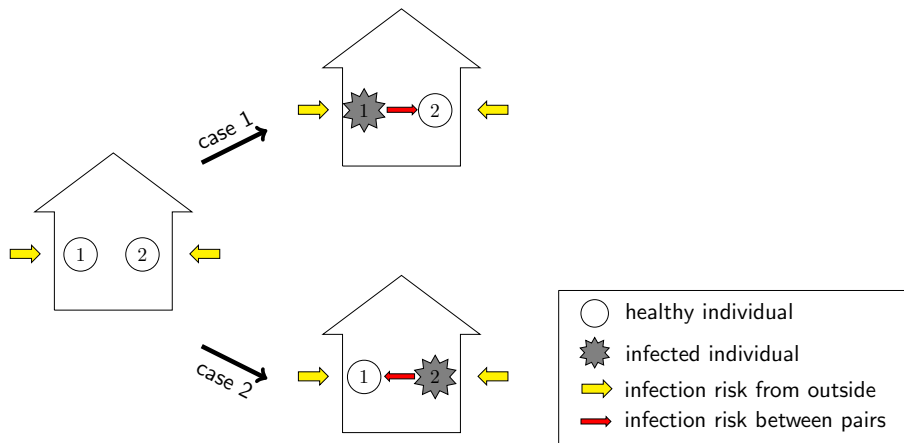
Consider two individuals with treatment  $X_1$  and  $X_2$  and infection outcome  $Y_1$  and  $Y_2$ .



- The graph is not an acyclic directed graph (DAG).

## Two possibilities of the process

Separate the process into two distinctive possibilities: case 1 and case 2.



# Unique challenges for causal identification under contagion

Problem: Differentiating exposure to infection, even after randomization

## Solution:

Add a component of "exposure to infection" (other's infection times) into the counterfactual outcome definition for a fair comparison.

Problem: Bidirectional arrow in the causal diagram

## Solution:

Remove the bidirectional arrow by breaking down the transmission process into exclusive possibilities.

## Notation

For the symmetric partnership models, consider individual 1 and 2 and let,

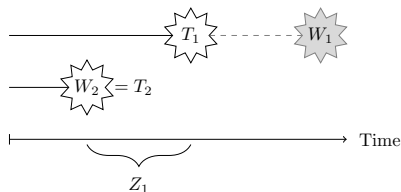
- Treatment assignment:  $X = (X_1, X_2)$
- Infection time:  $T_i$  for  $i = 1, 2$
- Infection outcome:  $Y_i(t) \equiv \mathbb{1}\{t \geq T_i\}$  for  $i = 1, 2$

Instead of using the same  $T_i$  for both cases, introduce additional variables to distinguish two cases.

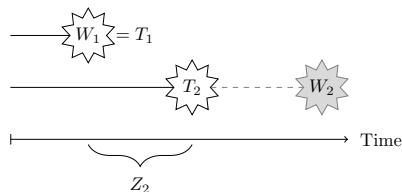
- Isolated infection time:  $W_i$  for  $i = 1, 2$
- Extra infection time after partner's infection:  $Z_i = T_i - W_j$  for  $i \neq j$



## Structure: Relationships between $W_i$ , $Z_i$ and $T_i$



Case 1: subject 2 gets infected first  
 $T_1, Z_1$  and  $W_2 = T_2$  observed  
 $W_1$  censored,  $Z_2$  undefined

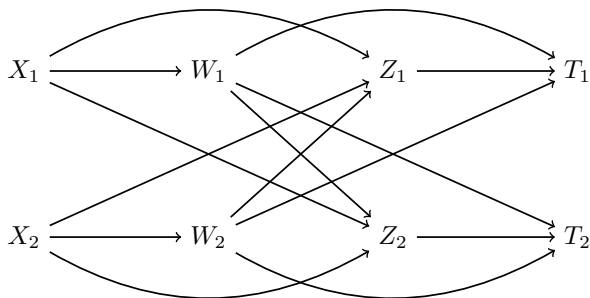


Case 2: subject 1 gets infected first  
 $T_2, Z_2$  and  $W_1 = T_1$  observed  
 $W_2$  censored,  $Z_1$  undefined

- $W_i$ : spontaneous infection time only by external risk
- $Z_i$ : additional time to infection after their partner's infection
- $T_i$ : infection time of  $i$

$$T_i = \begin{cases} W_i & \text{if } W_i < W_j \\ W_j + Z_i & \text{otherwise} \end{cases}$$

## Graphical representation

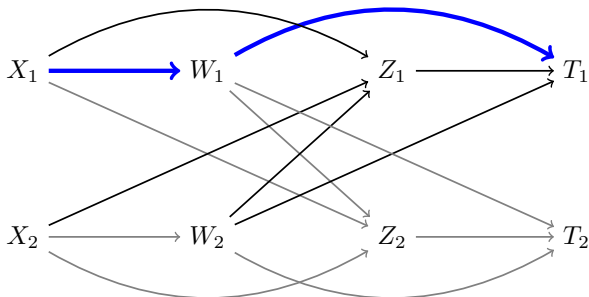


(Covariates  $L$  are omitted for simplicity)

- $W_i$ : spontaneous infection time only by external risk
- $Z_i$ : additional time to infection after their partner's infection
- $T_i$ : infection time of  $i$

## Graphical representation

Use individual 1 as an example,

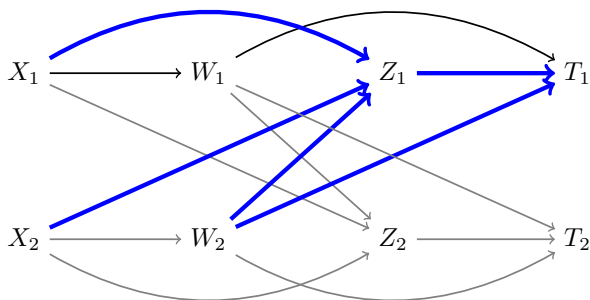


(Covariates  $L$  are omitted for simplicity)

- $W_i$ : spontaneous infection time only by external risk
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## Graphical representation

Use individual 1 as an example,



(Covariates  $L$  are omitted for simplicity)

- $W_i$ : spontaneous infection time only by external risk
- $Z_i$ : additional time to infection after their partner's infection
- $T_i$ : infection time of  $i$

## Notation

Define counterfactual infection outcome  $Y_i(t; s, x_i, x_j)$  for  $i = 1, 2$  where  $j \neq i$ , when we fix (i) its **own treatment**  $X_i = x_i$ , (ii) **partner's treatment**  $X_j = x_j$ , and (iii) **partner's infection time**  $W_j = s$ .

Note: partner's infection time is treated as another type of intervention for the potential outcomes – "exposure to infection".

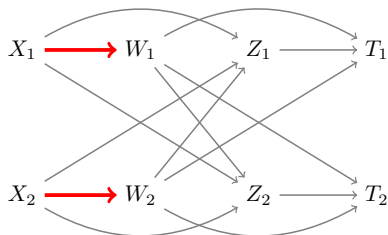
### Goal:

Identify  $Y_i(t; s, x_i, x_j)$ , or equivalently  $T_i(s, x_i, x_j)$ , under the joint intervention  $(s, x_i, x_j)$  for  $i = 1, 2$ .

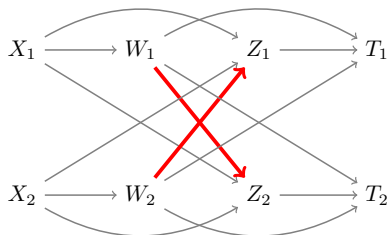
$$T_i(s, x_i, x_j) = \begin{cases} W_i(x_i) & \text{if } W_i(x_i) < W_j(x_j) \\ W_j(x_j) + Z_i(w_j(x_j); x_i, x_j) & \text{otherwise} \end{cases}$$

# Main Result: Assumptions

- Treatment exchangeability:  $Y_i(t; s, x) \perp\!\!\!\perp X \mid L$
- Consistency:  $Y_i(t) = Y_i(t; s, x)$  when  $X = x$  and  $W_j = s, j \neq i$
- Positivity:  $P(W_i = s, X = x | L = l) \in (0, 1)$  for all  $x \in \mathcal{X}, l \in \mathcal{L}$  and  $w_i > 0$  for  $i = 1, 2$
- Infection independence:  $W_i(x_i) \perp\!\!\!\perp W_j(x_j) | X, L$  for  $i \neq j$
- Infection ignorability:  $Z_i(t; s, x) \perp\!\!\!\perp W_j(x_j) | X, L$  for  $i \neq j$



Infection independence



Infection ignorability

# Main Result: Identification

## Identification theorem

Suppose Assumptions 1-5 hold. For fixed value of  $s$ ,  $t$  and  $\mathbf{x} = (x_i, x_j)$ , the average risk of infection by time  $t$  for the focal individual  $i$ , which is  $\mathbb{E}[Y_i(t; s, \mathbf{x})]$ , is identified as:

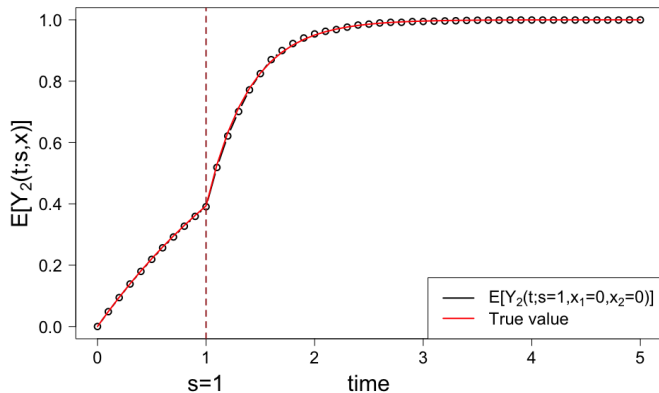
$$\mathbb{E}[Y_i(t; s, \mathbf{x})] = 1 \cdot p_i(s|\mathbf{x}) + \mathbb{E}[Y_i(t) | T_i \geq s, T_j = s, \mathbf{X} = \mathbf{x}] \cdot [1 - p_i(s|\mathbf{x})]$$

where  $p_i(s|\mathbf{x}) = 1 - \exp\left[-\int_0^s \frac{\Pr(T_i=u, T_j>u|\mathbf{X}=\mathbf{x})}{\Pr(T_i>u, T_j>u|\mathbf{X}=\mathbf{x})} du\right]$

Note: Instead of using binary infection outcome by the end of observation, this causal identification is built on observation of infection time, which provides sufficient control for exposure to infection.

## Simulation: causal identification for potential outcome

We simulate  $N=100,000$  partnerships with constant exogenous hazard 0.5 and within-pair hazard 2. Vaccinations both decrease risks for vaccinated and reduce transmissibility by 50%.





## Causal estimands: Exposure-controlled causal estimands

### Exposure-controlled causal estimands

- Susceptibility effect ( $s > 0$ )

$$SE(t, s, x_1) = \mathbb{E}[Y_i(t; s, \mathbf{1}, x_j) - Y_i(t; s, \mathbf{0}, x_j)]$$

- Infectiousness effect ( $s > 0$ )

$$IE(t, s, x_i) = \mathbb{E}[Y_i(t; s, x_i, \mathbf{1}) - Y_i(t; s, x_i, \mathbf{0})]$$

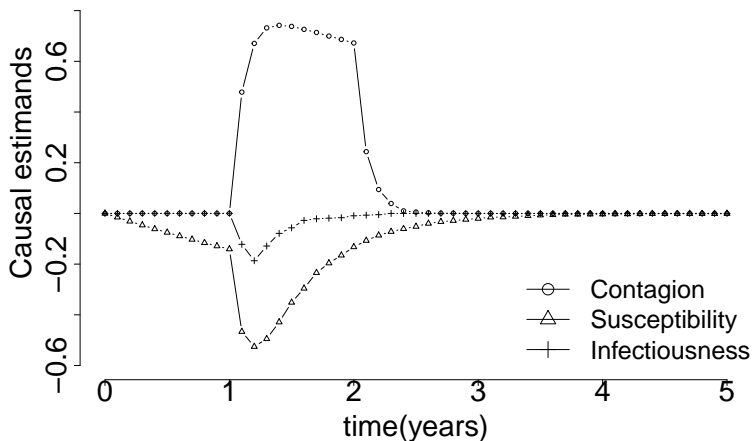
- Contagion effect ( $s \neq s'$  and  $X = (0, 0)$ )

$$CE(t, s, s') = \mathbb{E}[Y_i(t; s', 0, 0) - Y_i(t; s, 0, 0)]$$

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

## Simulation: Estimates of controlled causal estimands

We simulate  $N=100,000$  partnerships with constant exogenous hazard 0.5 and within-pair hazard 2. Vaccinations decrease infection risk of treated individuals and transmission ability both by 50%. We choose  $s = 1$  and  $s' = 2$ .



# Causal estimands: Exposure-marginalized causal estimands

## Identification for natural potential outcomes

$$\mathbb{E}[Y_i(t; W_j(x'_j), x)|L = 1] = \int_0^t \mathbb{E}[Y_i(t; w_j, x)|L = 1]dF_j(w_j|x'_j, l_j).$$

## Exposure-marginalized (natural) causal estimands

- Susceptibility effect

$$SE_i(t, x_j) = \mathbb{E}[Y_i(t; W_j(x_j), x_i = 1, x_j) - Y_i(t; W_j(x_j), x_i = 0, x_j)]$$

- Infectiousness effect

$$IE_i(t, x_i) = \mathbb{E}[Y_i(t; W_j(0), x_i, x_j = 1) - Y_i(t; W_j(0), x_i, x_j = 0)]$$

- Contagion effect

$$CE_i(t, x) = \mathbb{E}[Y_i(t; W_j(0), x_i, x_j) - Y_i(t; W_j(1), x_i, x_j)]$$

Other commonly used estimators:

- Direct effect:  $DE(t) = \mathbb{E}[Y_i(t)|X_i = 1] - \mathbb{E}[Y_i(t)|X_i = 0]$
- Indirect effect:  $IDE(t) = \mathbb{E}[Y_i(t)|X_j = 1] - \mathbb{E}[Y_i(t)|X_j = 0]$

# Simulation: Estimations of natural causal estimands

Disease transmission dynamic:

- Constant external and internal hazards
- time-varying external and internal hazards

Treatment assignment:

- Bernoulli, Block, and Cluster randomization
- Observational studies with correlated  $(X_i, X_j)$  due to confounders

Natural causal estimands to be compared:

- Exposure-marginalized susceptibility effect  $SE(t, 0)$
- Exposure-marginalized infectiousness effect  $IE(t, 0)$
- Exposure-marginalized contagion effect  $CE(t, 0, 0)$
- Direct effect  $DE(t)$
- Indirect effect  $IDE(t)$
- ... (others omitted, see details in paper)

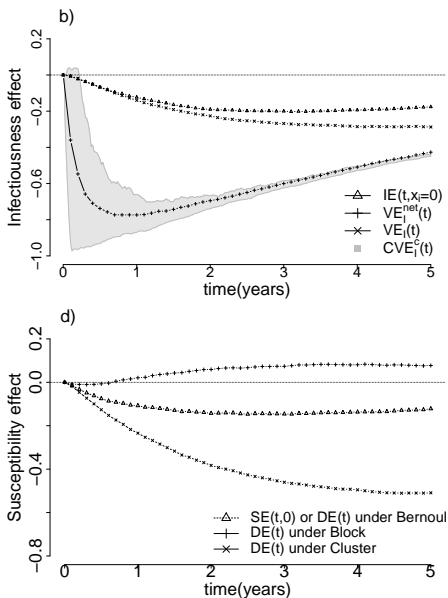
# Simulation: Estimations of natural causal estimands

Simulation	Treatment	$CE(t, 0, 0)$	$SE(t, 0)$	$IE(t, 0)$	DE(t)	IDE(t)
Constant hazards	Obs.	0.12	-0.14	-0.19	-0.16	-0.20
	Bernoulli	0.12	-0.14	-0.19	-0.16	-0.20
	Block	-	-	-	0.06	-
	Cluster	-	-	-	-0.39	-
Constant hazards without contagion	Obs.	0.00	-0.18	0.00	-0.18	0.00
	Bernoulli	0.00	-0.18	0.00	-0.18	0.00
	Block	-	-	-	-0.18	-
	Cluster	-	-	-	-0.18	-
Time-varying hazards	Obs.	0.12	-0.14	-0.20	-0.21	-0.22
	Bernoulli	0.12	-0.14	-0.20	-0.21	-0.22
	Block	-	-	-	0.08	-
	Cluster	-	-	-	-0.50	-
Time-varying hazards without contagion	Obs.	0.00	-0.28	0.00	-0.28	0.00
	Bernoulli	0.00	-0.28	0.00	-0.28	0.00
	Block	-	-	-	-0.28	-
	Cluster	-	-	-	-0.28	-

[1] VanderWeele et al. Effect partitioning under interference in two-stage randomized vaccine trials. *Statistics probability letters*, 81(7): 861-869, 2011.

[2] Eck et al. Randomization for the susceptibility effect of an infectious disease intervention. Submitted, 2019.

# Simulation: Estimations of natural causal estimands



# Summary

- we propose a new framework to articulate causal structure of infectious disease outcomes and treatments, in the case of partnership models.
- A class of fundamental (controlled- and marginalized-) causal estimands for the susceptibility, infectiousness and contagion effect of vaccines are proposed and identified non-parametrically.
- Comprehensive comparisons of our casual estimands are made to other popular estimands in contemporary epidemiology.

## Takehome message

- Causal estimands involving interactions with others (contagion and infectiousness effects) require control on others' infection times.
- Valid causal estimands depend on proper randomization scheme and proper control of an comparable distribution of the partner's infection time.
- Causal estimands are functions of observation time, underlying transmission dynamic, cluster size and many other factors.
- Direct comparison of causal estimands from different trails with different observational time and other features may not be meaningful.



## Further questions

- ① How do we generalize the identification technique to bigger clusters?
- ② How do we derive causal estimands, which do not depend on observation time, cluster size, transmission dynamic, or interactions between individuals?
- ③ How do we calculate causal estimands more efficiently for bigger clusters.

Welcome to the causal inference reading group at  
2:00 - 3:30 pm on December 1st, 2020 for further discussions about:  
"Causal identification of infectious disease intervention effects in a clustered population." Xiaoxuan Cai, Eben Kenah, and Forrest W. Crawford.

# Acknowledgement

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

<https://arxiv.org/abs/1912.04151>

## Cox-type hazard model for pairwise infection times

Assume previously infected individuals, along with a exogenous source of infection, impose **independent** and **competing** risks of disease transmission to the remaining uninfected individuals.

For all  $i$  and all infected individual  $j$ ,  $j \neq i$ , we consider a Cox-type hazard model for the pairwise infection times:

- **External** source of infection:

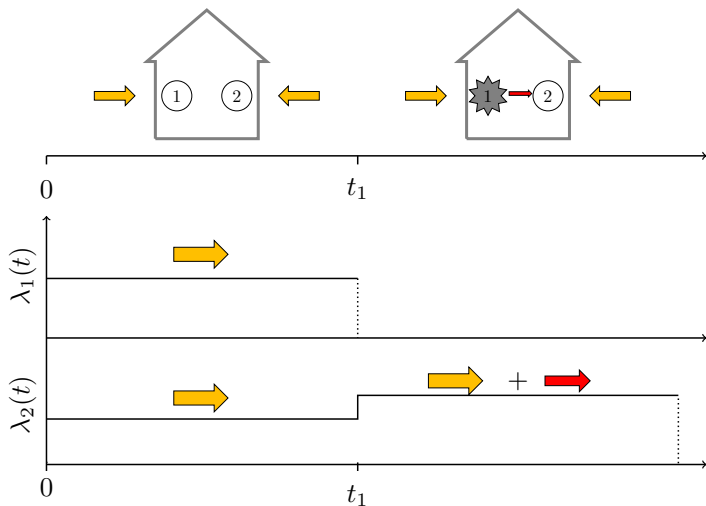
$$\lambda_{0i}(t | x_i, l_i) = \alpha(t) \exp[\beta_1 x_i + \theta_1^T l_i]$$

- **Internal** source of infection from infectious  $j$  to yet-uninfected  $i$ :

$$\lambda_{ji}(t | x_i, l_i) = \gamma(t - t_j) \exp[\beta_1 x_i + \beta_2 x_j + \theta_1^T l_i + \theta_2^T l_j]$$

where  $L = (L_i, L_j)$  measures baseline covariates for the two individuals, including shared covariates for the partnership as a whole.

# How do epidemiologists understand infectious disease transmission?



# Stochastic model for disease transmission

direct protection of vaccination

indirect protection of vaccination

$$\lambda_i(t) = \left( e^{\beta_1 x_i + \theta_1^T \mathbf{1}_i} \right) \times \left[ \underbrace{\alpha(t)}_{\text{time-varying hazard outside household}} + \sum_{j \neq i} \underbrace{y_j(t) \gamma(t - T_k)}_{\text{time-varying hazard from infectious household members}} e^{\beta_2 x_j + \theta_2^T \mathbf{1}_j} \right]$$

hazard<sub>i</sub>(t) = [susceptibility<sub>i</sub>] × [total exposure to infection(t)]

- $\alpha(t)$  is exogenous hazard of infection,  $\gamma(t)$  is endogenous hazard of transmission between individuals
- $\beta_1$  is for susceptibility effect,  $\beta_2$  is for infectiousness effect
- $\theta_1$  and  $\theta_2$  are covariate effects of susceptibility and infectiousness

# New vaccine estimands based on hazards

## Controlled hazard ratio vaccine effects

- Susceptibility hazard ratio:

$$HSE^C(t, s, x_j, l) = \frac{\lambda_i(t | s, 1, x_j, l)}{\lambda_i(t | s, 0, x_j, l)} = e^{\beta_1}$$

- Infectiousness hazard ratio:

$$HIE^C(t, s, s', x_i, l) = \frac{\lambda_i(t | s', x_i, 1, l) - \lambda_i(t | s, x_i, 1, l)}{\lambda_i(t | s', x_i, 0, l) - \lambda_i(t | s, x_i, 0, l)} = e^{\beta_2}$$

where  $s' < t < s$ .

- Contagion cumulative hazard ratio:

$$HCE^C(t; s'', s', l) = \frac{\int_0^t [\lambda_i(u; s', 0, 0, l) - \lambda_i(u; s, 0, 0, l)] du}{\int_0^t [\lambda_i(u; s'', 0, 0, l) - \lambda_i(u; s, 0, 0, l)] du} = \frac{\int_{s'}^s \gamma(u) du}{\int_{s''}^s \gamma(u) du}$$

where  $s' < s'' < t < s$ .

# Simulation: Estimations of causal estimands

Cluster	Treatment	Hazard estimands		Probability estimands				
		$\hat{\beta}_1$	$\hat{\beta}_2$	$\hat{CE}$	$\hat{SE}$	$\hat{IE}$	$DE(t)$	$IDE(t)$
Constant external and internal hazards								
2	Obs.	-0.91	-5.03	0.12	-0.14	-0.19	-0.16	-0.20
	Bernoulli	-0.90	-4.58	0.12	-0.14	-0.19	-0.16	-0.20
	Block	-0.88	-4.34	-	-	-	0.06	-
	Cluster	-0.91	-5.22	-	-	-	-0.39	-
Constant hazards without contagion								
2	Obs.	-0.91	-4.62	0.00	-0.18	0.00	-0.18	0.00
	Bernoulli	-0.89	-4.61	0.00	-0.18	0.00	-0.18	0.00
	Block	-0.88	-4.61	-	-	-	-0.18	-
	Cluster.	-0.92	-4.87	-	-	-	-0.18	-
Time-varying hazards								
2	Obs.	-0.88	-4.58	0.12	-0.14	-0.20	-0.21	-0.22
	Bernoulli	-0.93	-4.60	0.12	-0.14	-0.20	-0.21	-0.22
	Block	-0.87	-4.70	-	-	-	0.08	-
	Cluster	-0.92	-4.56	-	-	-	-0.50	-