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Abstract

When experimental subjects can interact with each other, the outcome of one individual may be affected by the treatment status of others. In many social science experiments, such spillover effects may occur through multiple networks, for example, through both online and offline face-to-face networks in a Twitter experiment. Thus, to understand how people use different networks, it is essential to estimate the spillover effect in each specific network separately. However, the unbiased estimation of these *network-specific spillover effects* requires an often-violated assumption that researchers observe all relevant networks. We show that, unlike conventional omitted variable bias, bias due to unobserved networks remains even when treatment assignment is randomized and when unobserved networks and a network of interest are independently generated. We then develop parametric and nonparametric sensitivity analysis methods, with which researchers can assess the potential influence of unobserved networks on causal findings. We illustrate the proposed methods with a simulation study based on a real-world Twitter network and an empirical application based on a network field experiment in China.

Keywords: causal inference, interference, potential outcomes, SUTVA, network experiments

1 Introduction

Existing methodologies for causal inference often assume the absence of spillover effects, that is, people are affected only by treatments directly assigned to them and not by those assigned to others (Cox 1958; Rubin 1980). However, in typical social science experiments, where individuals can interact with each other, spillover effects might naturally arise, and this causal interdependence across people is often of theoretical interest. Indeed, a growing number of political science studies use experiments to estimate spillover effects on a variety of outcomes, such as voting behavior (Nickerson 2008; Sinclair 2012; Sinclair *et al.* 2012; Foos and de Rooij 2017), electoral irregularities (Ichino and Schündeln 2012; Asunka *et al.* 2019; Bowers *et al.* 2018), the responsiveness of legislators (Coppock 2014), information diffusion in ethnic networks (Larson and Lewis 2017), and social norms in schools (Paluck *et al.* 2016).

In many of such social science applications, people can interact with each other through multiple channels, and thus, spillover effects often arise through multiple networks. For example, even though online network experiments typically focus only on online networks, people can also communicate with each other through their offline face-to-face network (Bond *et al.* 2012; Aral 2016; Coppock *et al.* 2016; Taylor and Eckles 2017). Other common types of multiple networks include friendship, neighbors, ethnic, kinship, and partisan networks, among many others (Fowler *et al.* 2011; Sinclair 2012). Therefore, it is important to estimate the spillover effect *specific* to each network. A substantive question is often not only about whether there exists any spillover effect but also about which networks people use to share information and influence each other's behavior. By estimating *network-specific spillover effects*, researchers can examine the mechanism of spillover effects.

However, the unbiased estimation of these network-specific spillover effects is challenging in practice. It requires making an often-violated assumption that researchers observe all relevant

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networks. For example, while scholars might carefully measure an ethnic network in a field experiment on ethnic voting, they might be unable to measure other types of network interactions, such as neighbors, friendship, and kinship networks. In this case, existing approaches, which assume no unobserved networks, can misattribute the spillover effects in unobserved networks to the ethnic network, resulting in biased estimates of network-specific spillover effects.

In this article, we address this methodological challenge by formally characterizing the bias due to unobserved networks and developing sensitivity analysis methods, with which researchers can assess the potential influence of unobserved networks on substantive findings.

In Section 2, we first extend the potential outcomes framework (Neyman 1923; Rubin 1974) to settings with multiple networks and then formally define the *average network-specific spillover effect* (ANSE). This new estimand represents the average causal effect of changing the treatment status of neighbors in a given network, without changing the treatment status of neighbors in other networks. The ANSE can be estimated without bias using an inverse probability weighting estimator as long as researchers can observe the network of interest and all other networks in which spillover effects exist. In Section 3, we then consider a scenario of unobserved networks and derive the exact bias formula. It is a function of the spillover effects through unobserved networks and the overlap between observed and unobserved networks. This bias formula implies that, unlike the conventional omitted variable bias, the bias for the ANSE is nonzero even when (a) treatment assignment is randomized and (b) unobserved networks and the network of interest are independently generated.

In Section 4, we propose parametric and nonparametric sensitivity analysis methods for evaluating the potential influence of unobserved networks on causal conclusions. Using these methods, researchers can derive simple formal conditions under which unobserved networks would explain away the estimated ANSE. Researchers can also use them to bound the ANSE using only two sensitivity parameters. Although the parametric sensitivity analysis method focuses on one unobserved network for the sake of intuitive interpretation, the nonparametric method can handle multiple unobserved networks.

We provide two empirical illustrations, each from the most popular types of network experiments; an online social network experiment and a network field experiment. The first is a simulation study based on the real-world Twitter network (Coppock *et al.* 2016) where we simulate a variety of unobserved offline face-to-face networks to assess the performance of the proposed approach (Section 5). The second is a reanalysis of the field network experiment in rural China (Cai *et al.* 2015). We apply the proposed sensitivity analysis methods and assess the robustness of the original findings to unobserved networks (Section 6).

Our paper builds on a growing literature on spillover effects (e.g., Hong and Raudenbush 2006; Sobel 2006; Rosenbaum 2007; Hudgens and Halloran 2008; Tchetgen Tchetgen and VanderWeele 2010), especially spillover effects in networks (e.g., Aronow 2012; Bowers *et al.* 2013; Toulis and Kao 2013; Liu and Hudgens 2014; Ogburn and VanderWeele 2014; Athey *et al.* 2018; Forastiere *et al.* 2016; Aronow and Samii 2017; Eckles *et al.* 2017; Tchetgen Tchetgen *et al.* 2017; Bowers *et al.* 2018). See Halloran and Hudgens (2016) for a recent review about spillover effects in general. The vast majority of the work has mainly focused on the case where all relevant networks are observed. Only recently has the literature begun to study the consequence of unobserved networks. One way to handle unobserved networks is to consider the problem as “misspecification” of the spillover structure (Aronow and Samii 2017; Sävje 2019). Although proposition 8.1 of Aronow and Samii (2017) implies that the inverse probability weighting estimator is biased for the ANSE unless all relevant networks are observed, the exact expression of bias is difficult to characterize in general settings. In this paper, we instead focus on one common type of misspecification—the network

of interest is observed, but other relevant networks are unobserved. We, therefore, can explicitly derive the exact bias formula and develop sensitivity analysis methods.

Other approaches to deal with unobserved networks include randomization tests (Luo *et al.* 2012; Rosenbaum 2007) and the use of a monotonicity assumption (Choi 2017). While these approaches are robust to unmeasured networks, estimands studied in these papers are designed to detect the total amount of spillover effects in all networks, rather than spillover effects specific to a particular network, which are the main focus of this paper. Sävje *et al.* (2017) discuss the estimation of the expected average treatment effect in the presence of unknown interference. While their result applies to general types of interference, their causal estimand is different from ours, that is, the direct effect of treatments rather than the spillover effect. Bhattacharya *et al.* (2019) propose to estimate underlying networks by structural learning algorithms under the assumption of a chain graph model. We take a different approach of the potential outcomes framework, and our focus is to characterize the exact bias and develop sensitivity analysis methods rather than recovering underlying networks.

2 Spillover Effects in Multiple Networks

Causal inference in randomized experiments often assumes no interference (Cox 1958; Rubin 1980), that is, units are affected only by treatments directly assigned to them, and not by treatments assigned to other units. However, when units are connected in networks, treatments can have spillover effects on other units, and this causal interdependence is of theoretical interest. While the recent literature on interference has focused on settings with one network, this section extends the potential outcomes framework to settings with multiple networks. This setup serves as the foundation for analyzing unobserved networks and developing methodologies in Sections 3 and 4.

As our running example, we consider an online social network experiment (Aral 2016; Taylor and Eckles 2017), such as political mobilization experiments on Facebook (e.g., Bond *et al.* 2012) and on Twitter (e.g., Coppock *et al.* 2016). Although typical online network experiments only measure online networks of interest, people are often embedded in other networks, most importantly, an offline face-to-face network. Thus, experimental subjects can share information with one another through Facebook as well as through their offline face-to-face interactions, inducing spillover effects in both online and offline networks. To introduce the multiple network framework, we focus on such spillover effects in online and offline face-to-face networks as an illustrative example. In Sections 5 and 6, we provide two detailed empirical illustrations; a simulation study based on the Twitter network (Coppock *et al.* 2016) and a reanalysis of the field network experiment in rural China (Cai *et al.* 2015), respectively.

2.1 The Setup

Consider a randomized experiment with sample size N and each unit is indexed by $i \in \{1, 2, \dots, N\}$. Define a treatment assignment vector $\mathbf{T} = (T_1, \dots, T_N)^\top$ where binary treatment variable $T_i \in \{0, 1\}$ denotes the treatment received by unit i . For example, $T_i = 1$ would mean that unit i receives an informational message, and $T_i = 0$ would mean unit i is in a control group who does not receive any message. Based on the experimental design, treatment assignment probability $\Pr(\mathbf{T} = \mathbf{t})$ is known for all $\mathbf{t} \in \{0, 1\}^N$. Using the potential outcomes framework (Neyman 1923; Rubin 1974), let $Y_i(\mathbf{t})$ denote the potential outcome of individual i if the treatment assignment vector \mathbf{T} is set to \mathbf{t} . Importantly, the potential outcome of individual i is affected not only by her own treatment assignment, but also by the treatments received by others. Thus, we allow for spillover/interference between individuals (Cox 1958; Rubin 1980). In our running example, voting behavior of a given individual can be affected not only by whether she directly receives the informational

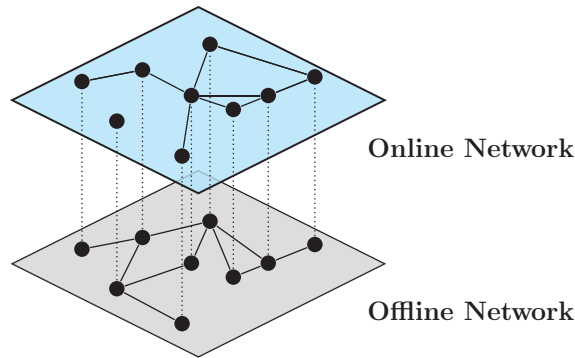


Figure 1. Example of two networks. *Note:* The same set of people are connected differently by an online network and an offline network. Neighbors in the two networks overlap, i.e., some, but not all, pairs of people are connected in both online and offline networks.

message, but also by messages assigned to her friends as people can share information with one another.

To formalize whose treatment status can affect a given individual, we rely on networks. In particular, consider two networks \mathcal{G} and \mathcal{U} connecting units with different edge sets. For example, people can be connected with Facebook and their offline face-to-face network. Formally, we define an adjacency matrix of network \mathcal{G} to be $A^{\mathcal{G}}$ where $A_{ij}^{\mathcal{G}} = A_{ji}^{\mathcal{G}} = 1$ if unit i is connected to unit j (and zeros on the diagonal). We can also incorporate directed ties when we define $A_{ij}^{\mathcal{G}} = 1$ if unit i “follows” unit j , for example, on Twitter. By assigning different weights to each tie, this framework can accommodate the strength of ties as well. We then define individual i ’s neighbors in network \mathcal{G} to be other individuals to whom she is connected in network \mathcal{G} , formally, $\mathcal{N}_i^{\mathcal{G}} \equiv \{j : A_{ij}^{\mathcal{G}} \neq 0\}$. We define $A^{\mathcal{U}}$ and $\mathcal{N}_i^{\mathcal{U}}$ similarly. For example, neighbors are those with whom individual i is friends on Facebook ($\mathcal{N}_i^{\mathcal{G}}$; neighbors on an online network) or those who individual i meets in person on a daily basis ($\mathcal{N}_i^{\mathcal{U}}$; neighbors on an offline face-to-face network). Importantly, neighbors in the two networks can overlap; we meet some but not all of our Facebook friends on a daily basis. Figure 1 visualizes an example with two networks.

We explicitly incorporate these two networks into the potential outcomes. In particular, we extend the stratified interference assumption (Hudgens and Halloran 2008) to multiple networks. We assume that the potential outcomes of individual i are affected by her own treatment assignment and the treated proportions of neighbors in networks \mathcal{G} and \mathcal{U} (Manski 2013; Toulis and Kao 2013; Liu and Hudgens 2014; Forastiere et al. 2016).

Assumption 1 (Stratified Interference with Multiple Networks) For all i ,

$$Y_i(\mathbf{t}) = Y_i(d, \mathbf{g}, \mathbf{u}),$$

where d is her own treatment assignment, \mathbf{g} and \mathbf{u} are the proportions of treated neighbors in networks \mathcal{G} and \mathcal{U} , respectively:

$$d = t_i, \quad \mathbf{g} = \frac{\sum_{j \in \mathcal{N}_i^{\mathcal{G}}} t_j}{|\mathcal{N}_i^{\mathcal{G}}|}, \quad \mathbf{u} = \frac{\sum_{j \in \mathcal{N}_i^{\mathcal{U}}} t_j}{|\mathcal{N}_i^{\mathcal{U}}|},$$

where t_i is the treatment received by individual i . $\sum_{j \in \mathcal{N}_i^{\mathcal{G}}} t_j$ is the total number of treated units among individual i ’s neighbors in network \mathcal{G} and $|\mathcal{N}_i^{\mathcal{G}}|$ is the total number of individual i ’s neighbors in network \mathcal{G} . $\sum_{j \in \mathcal{N}_i^{\mathcal{U}}} t_j$ and $|\mathcal{N}_i^{\mathcal{U}}|$ are similarly defined.

The potential outcomes depend on the treatment assignment of herself d and the fractions of treated neighbors in each network (g, u) . For example, the potential outcome of unit i is a function of whether she receives the informational message (d), the proportion of Facebook friends who receive the message (g), and the proportion of offline friends who receive the message (u). Although this assumption can only allow for spillover effects through treated proportions in multiple networks, it is more flexible than the conventional assumption of no interference, which requires that the treatment status of neighbors do not change potential outcomes at all. Formally, Assumption 1 can also be viewed as an exposure mapping $f(\mathbf{t}, (A_i^{\mathcal{G}}, A_i^{\mathcal{U}}))$ set to a three-dimensional vector (d, g, u) (Aronow and Samii 2017).

Several points about Assumption 1 are worth clarifying. First, because this assumption is made at the individual level, an equivalent statement of Assumption 1 is that the potential outcomes depend on the treatment assignment of herself and the number (rather than proportions) of treated neighbors in each network. The only change is that when defining causal estimands, we should explicitly condition on the total number of neighbors in each network. Second, Assumption 1 is violated when neighbors in the same network have different spillover effects. For example, suppose a given individual i has three Facebook friends $\{F_1, F_2, F_3\}$, and F_1 has a larger spillover effect than the other two. In this case, even when $g = 1/3$, the potential outcomes of unit i differ depending on whether F_1 is treated or one of the other two is treated, which violates Assumption 1. Third and most importantly, unlike the previous literature, Assumption 1 is defined with multiple networks. Therefore, researchers can make the assumption more plausible by specifying networks more precisely. In the example above, suppose F_1 has a larger spillover effect than the other two because F_1 is also a friend of unit i on Twitter, whereas the other two are not. Then, we can restore Assumption 1 with respect to three networks; the Facebook network, the Twitter network, and an offline network. However, it is important to note that as we consider more networks, it is more difficult to observe all relevant networks, which is the central topic of the paper we further discuss in Sections 3 and 4.

We define networks to be *causally relevant* if treated proportions in such networks have nonzero causal effects.

Definition 1 (Relevant Networks) Network \mathcal{G} is causally relevant if $Y_i(d, g, u) \neq Y_i(d, g', u)$ for some i and $(d, g, u), (d, g', u) \in \Delta_i$ where Δ_i is the support of treatment exposure vector (T_i, G_i, U_i) . The relevance of network \mathcal{U} is similarly defined.

For example, if people do not talk about elections at all on Facebook, Facebook would be causally irrelevant because messages received by other Facebook friends would have no effect on voting behavior through Facebook when fixing treated proportions in the offline network.

Finally, as in the standard causal inference setting, we observe only one of many potential outcomes. For all i , $Y_i = \sum_{(d, g, u) \in \Delta_i} 1\{T_i = d, G_i = g, U_i = u\} Y_i(d, g, u)$ (Holland 1986).

2.2 Causal Quantities of Interest

Without loss of generality, we assume two networks \mathcal{G} and \mathcal{U} are causally relevant and define our quantities of interest using these two networks.

- 2.2.1 *Direct Effects and Network-Specific Spillover Effects.* First, we define the direct effect of a treatment. It is the difference between the potential outcomes under treatment and control, averaging over the distribution of treatment assignment (G_i, U_i) . Formally, we define the *average direct effect* (ADE) as follows.

Definition 2 (Average Direct Effect)

$$\delta \equiv \frac{1}{N} \sum_{i=1}^N \left\{ \sum_{(g,u) \in \Delta_i^{gu}} \{Y_i(1, g, u) - Y_i(0, g, u)\} \Pr(G_i = g, U_i = u) \right\}, \quad (1)$$

where the support is defined as $\Delta_i^{gu} = \{(g, u) : \Pr(G_i = g, U_i = u) > 0\}$.

It is the weighted average of the causal effects, $Y_i(1, g, u) - Y_i(0, g, u)$, which hold the proportions of treated neighbors in the two networks constant. Intuitively, this effect quantifies the causal impact of the treatment directly received by herself. Thus, the ADE could represent the direct causal effect of the informational message on voting behavior. The ADE is a simple extension of the expected average treatment effect (Sävje *et al.* 2017) to the two-network settings. We keep the term of the *direct* effect in order to distinguish it from spillover effects we define next.

Spillover effects describe the causal effects of the neighbors' treatment status on a given individual. It is the change in the potential outcome when the proportion of treated neighbors goes from a lower proportion to a higher proportion. Formally, it is the difference between the potential outcome for a given individual when $g^H \times 100\%$ of her neighbors in \mathcal{G} are treated and the other potential outcome for the same individual when $g^L \times 100\%$ of her neighbors in \mathcal{G} are treated, holding her own treatment assignment and averaging over the proportions of treated neighbors in \mathcal{U} . Two constants g^H and g^L stand for "Higher" and "Lower" percent. We define the *average network-specific effect* (ANSE) as follows.

Definition 3 (Average Network-Specific Spillover Effect)

$$\tau(g^H, g^L; d) \equiv \frac{1}{N} \sum_{i=1}^N \left\{ \sum_{u \in \Delta_i^u} \{Y_i(d, g^H, u) - Y_i(d, g^L, u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L) \right\}, \quad (2)$$

where the support is defined as $\Delta_i^u = \{u : \Pr(U_i = u \mid T_i = d, G_i = g^H) > 0 \text{ and } \Pr(U_i = u \mid T_i = d, G_i = g^L) > 0\}$. The ANSE for network \mathcal{U} is defined similarly.

It is the weighted average of the spillover effects specific to network \mathcal{G} , $Y_i(d, g^H, u) - Y_i(d, g^L, u)$, which hold her own treatment assignment and the proportion of treated neighbors in \mathcal{U} constant. Thus, for example, the ANSE ($g^H = 0.8, g^L = 0.2; d = 0$) could represent the Facebook-specific spillover effect where we change the treated proportion of one's Facebook friends (from $g^L = 20\%$ to $g^H = 80\%$) while holding constant one's own treatment assignment ($d = 0$) and averaging over the treated proportion of one's offline friends (u). Similarly, the ANSE ($u^H = 0.8, u^L = 0.2; d = 0$) could represent the offline network-specific spillover effect where we change the treated proportion of one's offline friends (from $u^L = 20\%$ to $u^H = 80\%$) while holding constant one's own treatment assignment ($d = 0$) and averaging over the treated proportion of one's Facebook friends (u).

Importantly, the ANSE captures the spillover effect specific to each network separately. Thus, when Facebook (the offline network) is causally irrelevant, as defined in Definition 1, the ANSE in Facebook (the offline network) will be zero. By estimating the ANSE for each network, researchers can unpack the mechanism of spillover effects; which networks do people use to share information and influence each other's voting behavior? In Appendix B, we additionally show that the ANSE decomposes the total spillover effect, a popular estimand in the literature (Hudgens and Halloran 2008), into each network.

2.3 Estimation

In this section, we study the estimation of the ADE and the ANSE. We begin by introducing a common assumption made in practice; the *no omitted network* assumption defined below.

Assumption 2 (No Omitted Network) All causally relevant networks are observed.

For example, in the Facebook mobilization experiment (Bond *et al.* 2012), this assumption of no omitted network means that the Facebook network is the only relevant network and an unobserved face-to-face network is irrelevant; experimental subjects could affect one another through Facebook but not through their unobserved offline interactions. We first examine the estimation under this assumption and then discuss its violation in the subsequent sections.

Following the recent literature (e.g., Hudgens and Halloran 2008; Tchetgen Tchetgen and VanderWeele 2010; Kang and Imbens 2016; Sussman and Airoldi 2017), we start with design-based, inverse probability weighting (IPW) estimators for the ADE and ANSE (Horvitz and Thompson 1952; Aronow and Samii 2017). Importantly, researchers can compute the treatment exposure probability $\Pr(T_i = d, G_i = g, U_i = u)$ from the experimental design under the no omitted network assumption.¹ Thus, we rely on the following weighting estimators.

Theorem 1 (Estimation of the ADE and the ANSE) Under Assumption 2, the treatment exposure probability $\Pr(T_i = d, G_i = g, U_i = u)$ is known for all $(d, g, u) \in \Delta_i$ and all i . Therefore, the average direct effect and the average network-specific spillover effect in \mathcal{G} can be estimated by the following inverse probability weighting estimators.

$$\mathbb{E}[\hat{\delta}] = \delta \quad \text{and} \quad \mathbb{E}[\hat{\tau}(g^H, g^L; d)] = \tau(g^H, g^L; d),$$

where the expectation is taken over the experimental design $\Pr(\mathbf{T} = \mathbf{t})$, and

$$\hat{\delta} \equiv \frac{1}{N} \sum_{i=1}^N 1\{T_i = 1\} \tilde{w}_i Y_i - \frac{1}{N} \sum_{i=1}^N 1\{T_i = 0\} \tilde{w}_i Y_i \tag{3}$$

$$\hat{\tau}(g^H, g^L; d) \equiv \frac{1}{N} \sum_{i=1}^N 1\{T_i = d, G_i = g^H\} w_i Y_i - \frac{1}{N} \sum_{i=1}^N 1\{T_i = d, G_i = g^L\} w_i Y_i, \tag{4}$$

and weights are defined as

$$\tilde{w}_i = \frac{1}{\Pr(T_i | G_i, U_i)}, \quad \text{and} \quad w_i = \frac{\Pr(U_i | T_i, G_i = g^L)}{\Pr(U_i | T_i, G_i)} \times \frac{1}{\Pr(T_i, G_i)}.$$

We provide the proof in Appendix C.1. Although the IPW estimators are unbiased, researchers might want to increase efficiency by incorporating covariate information and some parametric assumptions (Särndal *et al.* 1992). In particular, we build on the design-based IPW estimator above and propose a weighted linear regression estimator. This alternative approach reduces standard errors at the expense of some bias due to parametric modeling assumptions. In particular, we consider

¹ It is important to emphasize that the treatment exposure probability is a function of both network structure and experimental design.

$$Y_i \sim \alpha + \beta T_i + \gamma_0 G_i + \gamma_1 (T_i \times G_i) + \mathbf{X}_i^\top \theta \text{ with weights } w_i. \tag{5}$$

The key assumptions are the linearity of G_i and the inclusion of pretreatment covariates \mathbf{X}_i .² We emphasize an important tradeoff. On the one hand, both assumptions are parametric in the sense that they are not directly justified by the experimental design. On the other hand, they tend to greatly reduce variance. Especially in settings with multiple networks, treatment exposure probabilities $\Pr(T_i, G_i, U_i)$ are usually small, and in many applied contexts, the standard IPW estimator has large standard errors. The weighted linear regression estimator aims to balance this bias-variance tradeoff in practice. See Toulis and Kao (2013) for a Bayesian approach, and Särndal *et al.* (1992); Rosenbaum (2002) and Aronow and Samii (2017) for the design-based covariate adjustment.

3 Exact Bias Formula

We can obtain an unbiased estimate of the ANSE when we observe all relevant networks. However, this assumption of no omitted network is often violated in practice, and if so, the ANSE even in the observed network cannot be estimated without bias. For example, in Bond *et al.* (2012), while the Facebook network is observed, an offline face-to-face network is unobserved. In this case, even with randomized experiments, an estimate of the Facebook-specific spillover effect is biased because people can potentially share information with offline friends. Bond *et al.* (2012) write, “it is plausible that unobserved face-to-face interactions account for at least some of the social influence that we observed in this experiment” (p. 298). In this section, to explicitly characterize sources of such bias, we derive the exact bias formula for the ANSE. We provide the exact bias formula for the ADE in Appendix C.2.1.

We consider a common research setting in which the main network of interest is observed but other relevant networks are not observed. In particular, we assume \mathcal{G} is an observed network of interest and \mathcal{U} is unobserved.³ Thus, the quantity of interest is the ANSE in the observed network \mathcal{G} . For simplicity, we refer to the ANSE in \mathcal{G} as the ANSE, without explicitly mentioning \mathcal{G} .

Since network \mathcal{U} is unmeasured, we cannot use weights w_i discussed in Section 2.3. Instead, we can only rely on partial weights $w_i^B \equiv 1/\Pr(T_i, G_i)$ where we only adjust for the direct treatment assignment T_i and the treated proportion in the observed network G_i . For example, the IPW estimator based on such partial weights becomes

$$\frac{1}{N} \sum_{i=1}^N \{1\{T_i = d, G_i = g^H\} w_i^B Y_i - 1\{T_i = d, G_i = g^L\} w_i^B Y_i\}. \tag{6}$$

We use $\hat{\tau}_B(g^H, g^L; d)$ to denote any estimator with partial weights w_i^B , including both the IPW estimator and the regression estimator discussed in Section 2.3. For the regression estimator, the bias formula we derive next can be seen as the lower bound where we focus only on the bias due to unobserved networks and not on bias due to functional form assumptions.

The next theorem shows the exact bias formula for $\hat{\tau}_B(g^H, g^L; d)$ in settings where the no omitted network assumption does not hold.

2 It is also possible to incorporate higher-order polynomial terms of G_i .
 3 If even the main network of interest is partially unobserved, it is impossible to identify the ANSE without strong assumptions because the treatment variable itself (the proportion of treated neighbors in the main network) is unobserved. If researchers are interested in estimating the ANSE in the observed part of the main network, the same results in Sections 3 and 4 hold by viewing the unobserved part of the main network as a separate unobserved network.

Theorem 2 (Bias for the ANSE due to Omitted Networks) When the no omitted network assumption (Assumption 2) is violated, estimator $\hat{\tau}_B(\mathbf{g}^H, \mathbf{g}^L; d)$ is biased for the ANSE.

$$\begin{aligned} & \mathbb{E}[\hat{\tau}_B(\mathbf{g}^H, \mathbf{g}^L; d)] - \tau(\mathbf{g}^H, \mathbf{g}^L; d) \\ &= \frac{1}{N} \sum_{i=1}^N \left\{ \sum_{u \in \Delta^u} \{Y_i(d, \mathbf{g}^H, u) - Y_i(d, \mathbf{g}^H, u')\} \{ \Pr(U_i = u | T_i = d, G_i = \mathbf{g}^H) - \Pr(U_i = u | T_i = d, G_i = \mathbf{g}^L) \} \right\}, \end{aligned}$$

for any $u' \in \Delta^u$.

We provide the proof in Appendix C.2. This bias can be decomposed into two parts: (1) the spillover effects attributable to the unobserved network \mathcal{U} , $Y_i(d, \mathbf{g}^H, u) - Y_i(d, \mathbf{g}^H, u')$ and (2) the dependence between the fraction of treated neighbors in \mathcal{G} and the fraction of treated neighbors in \mathcal{U} , $\Pr(U_i = u | T_i = d, G_i = \mathbf{g}^H) - \Pr(U_i = u | T_i = d, G_i = \mathbf{g}^L)$.

Based on this decomposition, we offer several implications of the theorem. First, when treatment assignment to \mathcal{U} has no effect (i.e., \mathcal{U} is irrelevant), $Y_i(d, \mathbf{g}^H, u) - Y_i(d, \mathbf{g}^H, u') = 0$ for any i . In this simple case, the bias is zero; this formula includes the assumption of no omitted network as a special case.

Second, the dependence between the fraction of treated neighbors in \mathcal{G} and the fraction of treated neighbors in \mathcal{U} determines the size and sign of the bias. In theory, when G_i and U_i are independent given T_i , the bias is zero. However, G_i and U_i are in general dependent given T_i . Two points about this dependence are worth noting. First, some randomization of treatment assignment, such as a Bernoulli design, can make T_i independent of (G_i, U_i) , but G_i and U_i are dependent given T_i even after any randomization of treatment assignment because some neighbors in \mathcal{G} are also neighbors in the other network \mathcal{U} ; networks \mathcal{G} and \mathcal{U} overlap each other. Formally, G_i and U_i are dependent because both are functions of the treatment assignment to common neighbors. Second, based on the same logic, G_i and U_i are not independent given T_i even when two networks \mathcal{G} and \mathcal{U} are independently generated because the two networks can still overlap each other.

For example, the Facebook network and an unobserved face-to-face network are likely to overlap each other. For some users, their Facebook friends are also close friends to whom they have offline interactions and vice versa. As long as the spillover effects in the face-to-face network are non-zero, an estimator ignoring this unobserved offline network (Equation (6)) would be biased for the Facebook-specific spillover effect.

Finally, we illustrate the bias with a simple linear model, $Y_i(T_i, G_i, U_i) = \alpha + \beta T_i + \gamma G_i + \lambda U_i + \epsilon_i$ where ϵ_i is an error term. Under the model, the potential outcome of individual i depends on neighbors in both \mathcal{G} and \mathcal{U} , and such two network-specific spillover effects do not interact. In this simple setup, the bias can be simplified as follows.

$$\mathbb{E}[\hat{\tau}_B(\mathbf{g}^H, \mathbf{g}^L; d)] - \tau(\mathbf{g}^H, \mathbf{g}^L; d) = \lambda \times \frac{1}{N} \sum_{i=1}^N \{ \mathbb{E}[U_i | T_i = d, G_i = \mathbf{g}^H] - \mathbb{E}[U_i | T_i = d, G_i = \mathbf{g}^L] \}.$$

It is clear that the bias depends on the ANSE in the unobserved network \mathcal{U} (i.e., λ) and the association between U_i and G_i given T_i . This bias is not zero unless the unobserved network \mathcal{U} is irrelevant because $\mathbb{E}[U_i | T_i = d, G_i = \mathbf{g}^H] \neq \mathbb{E}[U_i | T_i = d, G_i = \mathbf{g}^L]$ in general.

4 Sensitivity Analysis

We address the potential violation of the no omitted network assumption by developing parametric and nonparametric sensitivity analysis methods for the ANSE.

4.1 Parametric Sensitivity Analysis

Although the exact bias formula in Theorem 2 does not make any assumption about the unobserved network \mathcal{U} , in order to use it in applied work, it requires specifying a large number of sensitivity parameters. To construct a practical parametric sensitivity analysis method, we rely on a simplifying parametric assumption. In particular, we consider the assumption that the network-specific spillover effect in an unobserved network is linear and additive (Sussman and Airoidi 2017).

Assumption 3 (Linear, Additive Network-Specific Spillover Effect in \mathcal{U})

$$\frac{1}{\sum_{i=1}^N 1\{S_i = s\}} \sum_{i:S_i=s} \{Y_i(d, g, u) - Y_i(d, g, u')\} = \lambda(u - u'),$$

with some coefficient λ for all $(d, g, u), (d, g, u') \in \Delta_s$ where Δ_s is the support of (d, g, u) for i with the neighbors' profile $S_i = s$. The neighbors' profile S is defined such that the probability of treatment exposure is the same for those who have the same neighbors' profile. Formally, for $i \neq j$, $\Pr(T_i = d, G_i = g, U_i = u) = \Pr(T_j = d, G_j = g, U_j = u)$ when $S_i = S_j$. For example, when the Bernoulli or completely randomized design is used, S_i is simply a vector of three values; the number of neighbors in \mathcal{G} and \mathcal{U} , and the number of neighbors common to the two networks.

While this linear additive assumption is strong, it is commonly used in the causal inference literature to derive intuitive, easy-to-use sensitivity analysis methods. For example, the widely used sensitivity analysis methods for unobserved confounders in observational studies rely on similar assumptions (VanderWeele and Arah 2011). We also consider nonparametric sensitivity analysis in the next section under weaker assumptions.

Under Assumption 3, the general bias formula becomes the multiplication of two terms: the network-specific spillover effect in \mathcal{U} , that is, λ , and the effect of G_i on U_i given T_i , that is, $\frac{1}{N} \sum_{i=1}^N \{E[U_i | T_i = d, G_i = g^H] - E[U_i | T_i = d, G_i = g^L]\}$. The following theorem shows a simplified bias formula for settings with sparse \mathcal{G} .

Theorem 3 (Parametric Sensitivity Analysis) Under Assumption 3, a bias formula is approximated as follows.

$$E[\hat{\tau}_B(g^H, g^L; d)] - \tau(g^H, g^L; d) \approx \lambda \times \pi_{GU} \times (g^H - g^L), \tag{7}$$

where λ is the ANSE in network \mathcal{U} and π_{GU} is the overlap, that is, the fraction of neighbors in \mathcal{U} who are also neighbors in \mathcal{G} . Formally, $\pi_{GU} \equiv \sum_{i=1}^N \{|\mathcal{N}_i^{(\mathcal{G}, \mathcal{U})}| / |\mathcal{N}_i^{\mathcal{U}}|\} / N$ where $|\mathcal{N}_i^{(\mathcal{G}, \mathcal{U})}|$ is the number of unit i 's neighbors common to the two networks. If an experiment uses a Bernoulli design, the approximation is exact regardless of the sparsity of network \mathcal{G} .

We provide the proof in Appendix C.3. This simplified bias formula offers several implications. First, the bias is small when π_{GU} is small, that is, the overlap of neighbors in \mathcal{G} and \mathcal{U} is small. Hence, the bias is close to zero when the network \mathcal{G} is sparse and neighbors in \mathcal{G} and \mathcal{U} are disjoint. Second, even if two networks \mathcal{G} and \mathcal{U} are independently generated, the bias is not zero because $\pi_{GU} \neq 0$. We derive a similar parametric bias formula for settings with nonsparse \mathcal{G} in Appendix C.3.

For example, the overlap between the Facebook network and the unobserved face-to-face network is defined to be the fraction of offline friends who are also friends on Facebook. When this overlap is large (small), we expect the bias for the Facebook-specific spillover effect to be large (small).

To use this formula for a sensitivity analysis, researchers need to specify two sensitivity parameters: the network-specific spillover effect in an unobserved network (i.e., λ) and the fraction of neighbors in \mathcal{U} who are also neighbors in \mathcal{G} (i.e., π_{GU}). Once these two parameters are specified, we can derive the bias. Subsequently, because the bias utilizes only sensitivity parameters and $(g^H - g^L)$, we can obtain a bias corrected estimate by subtracting this bias from $\hat{\tau}_B(g^H, g^L; d)$. A sensitivity analysis is to report the estimated ANSE under a range of plausible values of λ and π_{GU} where $0 < \pi_{GU} < 1$. Note that $\lambda = 0$ corresponds to the no omitted network assumption.

4.2 Nonparametric Sensitivity Analysis

Now, we provide a nonparametric sensitivity analysis that makes only one assumption of non-negative outcomes. While we introduce our method using a random variable U_i for simplicity, the same method can be applied to a random vector \mathbf{U}_i and accommodate multiple unobserved networks. The proposed method is an extension of a sensitivity analysis developed for observational studies with no spillover effect (Ding and VanderWeele 2016) to experimental settings where spillover effects in unobserved networks induce bias.

As the parametric sensitivity analysis, we use two sensitivity parameters: intuitively, the network-specific spillover effect in \mathcal{U} , and the association between U_i and G_i . To capture the network-specific spillover effect in \mathcal{U} , let $MR_{UY}(g, s) \equiv \max_u \sum_{i; S_i=s} Y_i(d, g, u) / \min_u \sum_{i; S_i=s} Y_i(d, g, u)$ denote the largest potential outcomes ratios of U_i on Y_i given $T_i = d, G_i = g$ and $S_i = s$. For notational simplicity, we drop subscript d whenever it is obvious from contexts. Then, we define $MR_{UY} = \max_{g \in \{g^H, g^L\}, s \in \mathcal{S}} MR_{UY}(g, s)$ as the largest potential outcomes ratio of U_i on Y_i over $g \in \{g^H, g^L\}$ and $s \in \mathcal{S}$. Thus, MR_{UY} quantifies the largest possible potential outcomes ratio of U_i on Y_i . This ratio captures the magnitude of the network-specific spillover effect in \mathcal{U} . When network \mathcal{U} is irrelevant, $MR_{UY} = 1$.

Furthermore, to quantify the association between G_i and U_i , we use $RR_{GU}(g, g', u, s) = \Pr(U_i = u \mid T_i = d, G_i = g) / \Pr(U_i = u \mid T_i = d, G_i = g')$ to denote the relative risks of G_i on U_i for all i with $S_i = s$. $RR_{GU} = \max_{(g, g') \in \{g^H, g^L\}, u \in \Delta^u, s \in \mathcal{S}} RR_{GU}(g, g', u, s)$ is the maximum of these relative risks. This risk ratio captures the association between G_i and U_i .

Using these ratios, we can derive an inequality that the ANSE in the observed network \mathcal{G} needs to satisfy as long as outcomes are non-negative. The next theorem shows that we can obtain the bound for the ANSE with two sensitivity parameters, MR_{UY} and RR_{GU} .

Theorem 4 (Bound on the ANSE) When outcomes are non-negative,

$$\mathbb{E} \left[\frac{\hat{m}(d, g^H)}{B} - B \hat{m}(d, g^L) \right] \leq \tau(g^H, g^L; d) \leq \mathbb{E} \left[B \hat{m}(d, g^H) - \frac{\hat{m}(d, g^L)}{B} \right],$$

where $B = (RR_{GU} \times MR_{UY}) / (RR_{GU} + MR_{UY} - 1)$ and $\hat{m}(d, g) = \frac{1}{N} \sum_{i=1}^N 1\{T_i = d, G_i = g\} w_i^B Y_i$ for $g \in \{g^H, g^L\}$.

We provide the proof in Appendix C.4. Note that $MR_{UY} = 1$ corresponds to the no omitted network assumption, and $\mathbb{E}[\hat{m}(d, g^H) - \hat{m}(d, g^L)] = \mathbb{E}[\hat{\tau}_B(g^H, g^L; d)] = \tau(g^H, g^L; d)$ under the assumption. B is an increasing function of both RR_{GU} and MR_{UY} , implying that the bound is wider when the network-specific spillover effect in the unobserved network \mathcal{U} is larger and the effect of G_i on the distribution of U_i is larger. In fact, the size of the bound is given by $(B - \frac{1}{B})\mathbb{E}[\hat{m}(d, g^H) + \hat{m}(d, g^L)]$. It is important to note that the bound is not location-invariant because we use mean ratios and risk ratios as sensitivity parameters (Ding and VanderWeele 2016). The bound is valid as far as outcomes are non-negative, but how informative it is can vary. To conduct the sensitivity analysis, one can compute the bound for a range of plausible values of MR_{UY} and RR_{GU} . Compared

to the parametric sensitivity analysis, MR_{UY} and RR_{GU} correspond to λ and π_{GU} , respectively. Finally, as in Section 2.3, we can also use a weighted linear regression estimator for $\hat{m}(d, g)$ instead of the IPW estimator.

5 Simulation Study: Twitter Mobilization Experiment

Based on the real-world Twitter network (Coppock *et al.* 2016), we conduct a simulation study where we generate a variety of unobserved offline face-to-face networks. We evaluate how well the sensitivity analysis methods estimate the Twitter-specific spillover effect.

We vary the simulation setup along two dimensions; (1) the overlap between the observed Twitter network and a simulated unobserved offline network and (2) the outcome data generating process—a linear additive model or an interactive model. With this design, we illustrate three key results we derived analytically. First, the bias increases when the overlap between the online network and the unobserved offline network increases (Theorem 2). Second, the parametric sensitivity analysis can recover unbiased estimates under a linear additive model but suffers from bias under an interactive model (Theorem 3). Finally, the nonparametric sensitivity analysis provides bounds for the ANSE under both models, but it has larger confidence intervals than the parametric version under the linear additive model (Theorem 4).

Background.

To make our simulation as realistic as possible, we rely on the real-world Twitter network studied in Coppock *et al.* (2016). The original authors conducted a political mobilization experiment over the Twitter network to estimate the effectiveness of online mobilization appeals. In particular, they sampled followers of the Twitter account of a nonprofit advocacy organization, the League of Conservation Voters (LCV), and measured a network adjacency matrix among them. Each node of the network is a Twitter user who follows the LCV's account and a directed edge exists from user i to user j when user i follows user j . Instead of artificial simulated networks, we use this real-world Twitter network as the basis of our simulations after preprocessing as described below.

Simulation design.

To make the comparison of sensitivity analysis methods clear, we preprocess the Twitter network of Coppock *et al.* (2016). First, to avoid the influence of outliers, we remove units that are above the 95th percentile of the distribution on the size of Twitter neighbors (also known as the out-degree distribution). To have well-defined treated proportions G_i (explain more below), we also remove those who follow less than five other units. The resulting network contains 2,430 units with the mean number of neighbors equal to 18.5. We simulate an unobserved offline network \mathcal{U} with four different values of the overlap $\pi_{GU} \in \{0.0, 0.2, 0.4, 0.6\}$ where we expect no bias in the case of $\pi_{GU} = 0$. We use the Bernoulli design to randomly assign treatments with probability 0.5.

We consider two outcome data generating processes. For the linear additive model, we generate potential outcomes for individual i by $Y_i(T_i, G_i, U_i) = 5 + 2T_i + G_i + 1.5U_i + \epsilon_i$, where ϵ_i is randomly drawn from a normal distribution with $(\mu, \sigma) = (0.0, 0.5)$. G_i and U_i denote treated proportions of neighbors in the observed Twitter network and unobserved treated proportions of neighbors in the offline network, respectively. For the interactive model, we use $Y_i(T_i, G_i, U_i) = 5 + 2T_i + G_i + 2T_i \times U_i + 2G_i \times U_i + 0.5U_i + \epsilon_i$. As the main causal estimand, we consider the ANSE, $\tau(0.6, 0.4; 0)$, where we compare $g^H = 0.6$ relative to $g^L = 0.4$ when $d = 0$ (to have $G_i = 0.6$ and $G_i = 0.4$ both well-defined, we focus on a subset of units who follows at least five units). In the linear additive model, the true ANSE is equal to 0.2, and in the interactive model, the true ANSE is approximately equal to 0.4 (the exact values change according to the overlap π_{GU}).

The offline network \mathcal{U} is unmeasured and thus, we can rely only on a linear regression $Y_i \sim \alpha + \beta T_i + \gamma_0 G_i + \gamma_1 (T_i \times G_i)$ with weights $w_i^B = 1/\Pr(T_i, G_i)$. An estimator, $0.2 \times \hat{\gamma}_0$, is biased for

$\tau(0.6, 0.4; 0)$ due to unobserved network \mathcal{U} . The parametric sensitivity analysis provides a bias-corrected estimate, $0.2 \times \hat{\gamma}_0 - 0.2 \times \lambda \times \pi_{GU}$, with two sensitivity parameters; the overlap π_{GU} and the spillover effect in the unobserved network λ .⁴ The nonparametric sensitivity analysis computes bounds with sensitivity parameters (MR_{UY}, RR_{GU}).⁵ We evaluate an estimator that ignores unobserved offline networks, the parametric sensitivity analysis, and the nonparametric sensitivity analysis with the 2,000 Monte Carlo simulations.

Results.

Figure 2 presents simulation results. The first column reports estimates, $0.2 \times \hat{\gamma}_0$, that ignore the unobserved network \mathcal{U} , with the 95% confidence intervals (CI). Both settings of the linear additive and interactive models illustrate that the estimator is biased unless the overlap π_{GU} is zero, and the bias increases as the overlap increases. The second column shows the results for the parametric sensitivity analysis with the 95% CI. In the linear additive model setup (the first row), the parametric sensitivity analysis recovers approximately unbiased estimates. In contrast, in the interactive model setup (the second row), it still suffers from bias as the linear additive assumption (Assumption 3) is violated. Finally, the third column reports bounds from the nonparametric sensitivity analysis (thick bars) and the 95% CI of the bounds (thin bars). When the overlap π_{GU} is zero, the bound converges to an unbiased point estimate of the ANSE. Importantly, the bounds cover the true ANSE in both linear additive and interactive model settings as Theorem 4 implies. It also reveals an important limitation; while the nonparametric bounds cover the true ANSE, they are much wider than the 95% CI of the parametric sensitivity analysis under the linear additive model. This is the case especially when the overlap between the main network of interest and the unobserved network is large.

6 Empirical Application: Network Field Experiment

Cai *et al.* (2015) are interested in understanding how farmers in rural China use social networks to make important financial decisions, that is, insurance take-up. They designed an experiment with China's largest insurance provider, the People's Insurance Company of China (PICC), to randomly assign households to different information sessions about the insurance. While they estimate the direct treatment effect as a first step, the primary focus of the original analysis is to estimate the spillover effect of such information on the insurance take-up.

As in many other network field experiments, they use a social network survey to ask experimental subjects to name their network connections. In particular, they asked household heads to name five close friends with whom "they most frequently discuss rice production or financial issues" (Cai *et al.* 2015, p. 88). Although this is a popular strategy, there are several reasons to be afraid of unobserved networks. First, as carefully noted in the original paper, almost all households listed the maximum number of friends (the average number of listed friends is 4.9 where respondents can only list up to 5 friends), which suggests that respondents could have listed more friends if other forms of network surveys were used (Larson and Lewis 2020). Second, networks among experimental subjects are dense; "the average path-length is about 2.67, which means that a household can be connected to any other household in the village by passing on average of two to three households" (Cai *et al.* 2015, p. 89). This suggests that there are many potential network connections through which experimental subjects can communicate with each other. For example, in such rural Chinese villages, a kinship network is of great importance (e.g., Xiong and Payne 2017).

- 4 $\pi_{GU} \in \{0.0, 0.2, 0.4, 0.6\}$. In the linear additive model case, $\lambda = 1.5$. In the interactive model case, while λ is not well-defined, we use the main effect of U_i (0.5) as an example and show that the parametric sensitivity analysis cannot remove bias due to the violation of the linear additive assumption.
- 5 Imputing the potential outcome model into the definition, $MR_{UY} = 1.16$ (linear additive model) and $MR_{UY} = 1.14$ (interactive model). Using the definition of the overlap, $RR_{GU} = (0.6 \times \pi_{GU} + 0.5 \times (1 - \pi_{GU})) / (0.4 \times \pi_{GU} + 0.5 \times (1 - \pi_{GU}))$ (both models).

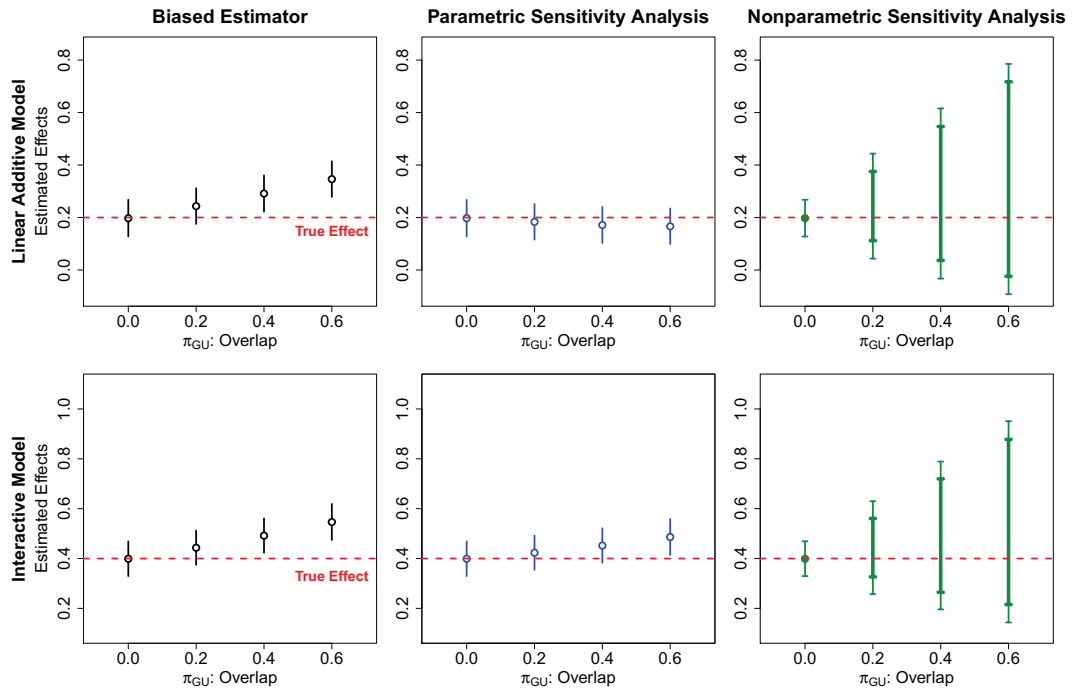


Figure 2. Simulation results on bias and sensitivity analysis for the Twitter experiment. *Note:* The first (second) row reports results from the linear additive (interactive) model. The first column demonstrates the bias in estimators that ignore the unobserved offline network. The second column shows that the parametric sensitivity analysis can recover unbiased estimates of the ANSE under the linear additive model but not under the interactive model. The third column demonstrates that bounds from the nonparametric sensitivity analysis cover the true ANSE, but are much wider than the parametric version under the linear additive model.

We extend the original analysis by estimating the spillover effect specific to the observed financial network while accounting for unobserved networks, such as the kinship network.

Setup.

Cai *et al.* (2015) designed the experiment with two rounds to estimate the spillover effect. A total of 2,175 households were invited to the first round and they were randomly assigned into one of two information sessions, *simple* or *intensive*. The *simple* session took about 20 min and the PICC agents explained the insurance contract. The *intensive* sessions took about 45 min and provided all the information given in the *simple* session, plus an additional detailed explanation of expected benefits and costs of purchasing the insurance. Three days after the first round, a different set of households were invited to the second round and randomly assigned into the *simple* or *intensive* sessions. We follow the original analysis and focus on 1,317 households who were invited to this second round and only received information from the *simple* or *intensive* session (657 and 660 households, respectively).⁶ The main outcome of interest is a binary variable whether each household in the second round takes up the insurance. Table 1 summarizes the relevant aspects of the experimental design. See Cai *et al.* (2015) for details and other features of the design.

For participants in the second round, the original authors define the main exposure variable of interest G_i to be the proportion of peers in their financial network who attended the first round *intensive* session. The direct treatment T_i is defined as an indicator variable taking 1 if household i receives the *intensive* session and 0 otherwise. Following their analysis, we focus on the ANSE

⁶ In the original paper, these groups are labeled as Simple2-NoInfo and Intensive2-NoInfo. There were two other randomly assigned groups who received additional information about the take-up rates in the first round. The original analysis (table 2 in the original paper) focuses on the first two groups, which we follow.

Table 1. Experimental design.

First round		Second round	
Simple	Intensive	Simple	Intensive
1079	1096	657	660
Total: 2,175		Total: 1,317	

specific to the observed financial network and compare $g^H = 0.2$ and $g^L = 0$ under no direct treatment receipt $d = 0$, formally $\tau(g^H = 0.2, g^L = 0; d = 0)$, as they show the biggest difference is between no treated peer and one treated peer (table 3 of the original paper). In addition, to satisfy the standard overlap assumption (Imbens and Rubin 2015), we analyze units who listed the maximum number of five households⁷ in the financial network question (1,207 households, more than 90% of all the samples).

To estimate spillover effects, the original authors estimate the following linear regression (model 4 in table 2 of Cai *et al.* (2015)) with standard errors clustered at the village level.

$$Y_i = \alpha + \beta T_i + \gamma_0 G_i + \gamma_1 (T_i \times G_i) + \mathbf{X}_i^\top \theta + \epsilon_i, \tag{8}$$

where pretreatment covariates \mathbf{X}_i include the size of neighbors, village fixed effects, and other household characteristics (gender, age, education of household head, rice production area, risk aversion, and perceived probability of future disaster). Thus, if the observed financial network is the only causally relevant network (i.e., the no omitted network assumption holds), an estimator $0.2 \times \hat{\gamma}_0$ is unbiased for the ANSE, $\tau(g^H = 0.2, g^L = 0; d = 0)$ where $\hat{\gamma}_0 = 0.444$ (SE = 0.109) in the original paper. However, as studied in Section 3, when the no omitted network assumption is violated—for example, people might share information through their kinship network, the estimator is biased for the ANSE specific to the financial network.

Sensitivity analysis.

We use the proposed sensitivity analysis methods to investigate the robustness of the original findings to unobserved networks. First, for the parametric sensitivity analysis, we consider three values for each of the two sensitivity parameters, the spillover effect in the unobserved network $\lambda \in (0.1, 0.2, 0.4)$ and the overlap $\pi_{GU} \in (0.2, 0.4, 0.6)$, producing a total of nine scenarios. The three values of λ are chosen to represent different scenarios in which we expect spillover effects in the kinship network is much smaller than, smaller than, or similar to the total spillover effect in the financial network ($\hat{\gamma}_0 = 0.444$). Similarly, the three values of π_{GU} cover from small to large overlaps. For the nonparametric sensitivity analysis, we also investigate three values for each of the two sensitivity parameters $MR_{UV} \in (1.3, 1.5, 1.8)$ and the overlap $RR_{GU} \in (1.3, 1.5, 1.8)$. Although these are relatively small risk and mean ratios (e.g., see table 1 in Ding and VanderWeele (2016)), we see below that substantive conclusions change according to different sensitivity parameters. Finally, while the original paper uses a linear regression without weights, we use a weighted linear regression with weights $w_i^B = 1/\Pr(T_i, G_i)$ as proposed in Section 4 for both analyses.

Figure 3 presents results from the parametric (left) and nonparametric sensitivity analysis (right). In the left panel, the leftmost point shows an estimate of the ANSE that ignores the unobserved kinship network with its 95% CI (8.87 percentage point; 95% CI = [3.31, 14.42]). The nine other points represent estimates from the parametric sensitivity analysis with their 95% CI. Importantly, point estimates are relatively stable over a range of sensitivity parameters

7 If the size of neighbors is smaller than five, $g^H = 0.2$ is impossible for such units, which violates the overlap assumption.

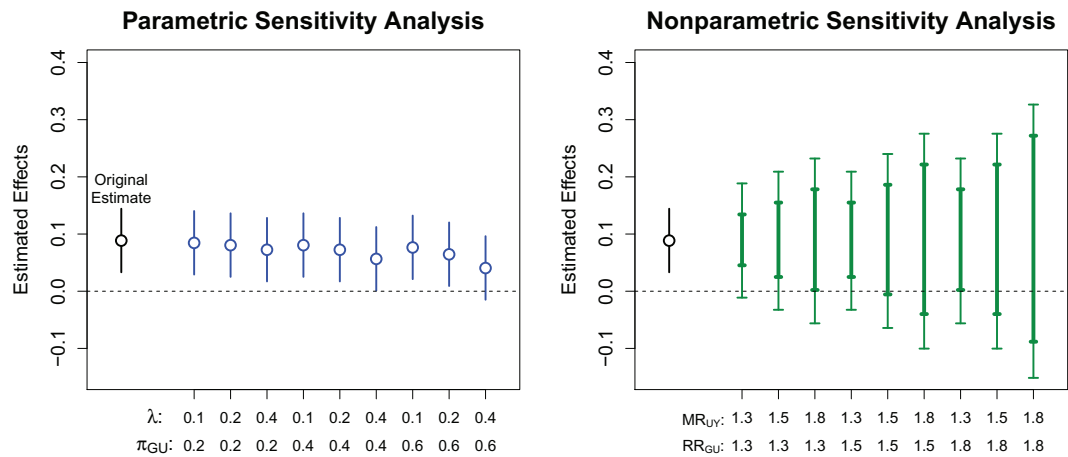


Figure 3. Parametric and nonparametric sensitivity analysis for the field network experiment in China. *Note:* The left (right) panel reports the parametric (nonparametric) sensitivity analysis with different sensitivity parameters. In both panels, the leftmost point shows an estimate of the ANSE that ignores unobserved networks. In the left panel, the nine other points represent estimates from the parametric sensitivity analysis with their 95% confidence intervals. In the right panel, the thick bars represent nonparametric bounds on the ANSE and the thin bars are their 95% confidence intervals.

and the 95% CI covers zero only when the effect in the unobserved kinship network is large ($\lambda = 0.4$) and the overlap between the observed financial network and the unobserved kinship network is relatively large ($\pi_{GU} = 0.6$). Thus, the parametric sensitivity analysis suggests that, while point estimates of the ANSE specific to the observed financial network are smaller than the original estimates, estimates are still positive and statistically significant at the conventional level of 0.05. Although this parametric sensitivity analysis is simple and intuitive, it requires the parametric linear additive assumption as discussed in Section 4.1. We now turn to the nonparametric sensitivity analysis which requires weaker assumptions at the expense of efficiency.

The right panel of Figure 3 shows results of the nonparametric sensitivity analysis where the leftmost point reproduces the same ANSE estimate that ignores the unobserved kinship network as a reference point. The thick bars represent nonparametric bounds on the ANSE and the thin bars are their 95% CI. Although lower bounds are positive for small bias scenarios, such as $(MR_{UY}, RR_{GU}) = (1.3, 1.3), (1.5, 1.3), (1.3, 1.5)$, all the 95% CI are inconclusive about the sign of the ANSE. Importantly, this result reveals that the positive ANSE estimates are sensitive to different assumptions about unobserved networks in the nonparametric sensitivity analysis, in contrast to the parametric sensitivity analysis above.

This difference in results from the parametric and nonparametric sensitivity analysis can arise for two reasons. First, the linear additive assumption (Assumption 3) required for the parametric sensitivity analysis could be violated and therefore, the nonparametric bounds are more credible. Second, the nonparametric bounds are in general less efficient than parametric methods and thus, large nonparametric bounds simply indicate insufficient statistical power for detecting the ANSE nonparametrically. In this China experiment, as the original authors detect the nonlinearity in the total spillover effect in the financial network (table 3 of Cai *et al.* (2015)), it is more likely that the linear additive assumption is violated and thus, the nonparametric sensitivity analysis is more credible. In practice, it is important to conduct both parametric and nonparametric sensitivity analysis methods as they are complementary and evaluate the robustness of the ANSE estimates under different assumptions.

7 Concluding Remarks

Although early work in the causal inference literature has assumed no interference, a growing number of both applied and methodological papers now explicitly incorporate spillover effects to understand causal interdependence across people. In this paper, we propose a framework for analyzing spillover effects in common social science settings of multiple networks, and introduce a new causal estimand, the average network-specific spillover effect (ANSE), to separately quantify the amount of spillover effects in each network. We show that the unbiased estimation of the ANSE requires an often-violated assumption of no omitted network. Unlike conventional omitted variable bias, the bias due to unobserved networks remains even when treatment assignment is randomized and when the network of interest and unobserved networks are independently generated. To account for this bias, we provide parametric and nonparametric sensitivity analysis methods, with which researchers can assess the robustness of causal conclusions to unobserved networks. The proposed methods are illustrated by two common types of network experiments; an online social network experiment and a network field experiment.

There are a number of future extensions that can further improve the methodologies proposed in this paper. First, although we made the assumption of stratified interference throughout this paper, we can potentially derive the exact bias formula and sensitivity analysis methods without it. This direction will be particularly important since not only the no omitted network assumption, but also the assumption of stratified interference might be strong in many applied settings. Second, while we focused on the estimation of the ANSE in this paper, it would also be of significant interest to study how to incorporate observed and unobserved networks into the rich literature of the Fisher randomization test for interference (e.g., Rosenbaum 2007; Aronow 2012; Bowers *et al.* 2013; Athey *et al.* 2018; Basse *et al.* 2019). Third, it is useful to study how we can effectively incorporate a supplementary survey to estimate the overlap between the main network of interest and an unobserved network. For example, in an online network experiment, even though it might be difficult to measure an offline face-to-face network for every subject in the experiment, researchers can use a network survey method (e.g., Bisbee and Larson 2017) to randomly sample subjects and use the estimated overlap for the sensitivity analysis. Fourth, this paper primarily focused on the bias due to unobserved networks, but another important extension would be to study implications of omitted relevant networks to variance estimation. A promising approach would be to consider a range of sensitivity parameters and compute the worst-case confidence interval (e.g., Berger and Boos 1994; Aronow *et al.* 2016). Finally, as discussed in the paper, the estimation accuracy is often of concern in settings with multiple networks. Thus, it would be useful to extend the literature on experimental design for spillover effects to settings of multiple networks and study how to design the optimal experiment (e.g., Hudgens and Halloran 2008; Sinclair *et al.* 2012; Basse and Airoidi 2018; Bowers *et al.* 2018; Jagadeesan *et al.* 2020).

Supplementary Appendix Appendix A. Details of Setup

We describe regularity conditions for the support of treatment exposure probabilities to ensure well-defined causal estimands.

The required regularity conditions are as follows: (1) the support of $\Pr(G_i = g, U_i = u | T_i = 1)$ is equal to the support of $\Pr(G_i = g, U_i = u | T_i = 0)$ for all i , and (2) the support of $\Pr(U_i = u | T_i = d, G_i = g^H)$ is equal to the support of $\Pr(U_i = u | T_i = d, G_i = g^L)$ for all i . We discuss them in order.

When we define the unit level direct effect, we avoid ill-defined causal effects by focusing on settings where the support of $\Pr(G_i = g, U_i = u | T_i = 1)$ is equal to the support of $\Pr(G_i = g, U_i = u | T_i = 0)$ for all i . This can be violated when the total number of treated units is small so that for some (g, u) , $\Pr(G = g, U = u | T_i = 1) = 0$ and $\Pr(G = g, U = u | T_i = 0) > 0$. One extreme example is that when we use complete randomization with the total number of treated units equal to 1. In this case,

whenever $T_i = 1, \Pr(G = g, U = u | T_i = 1) = 0$ for all (g, u) , but when $T_i = 0, \Pr(G = g, U = u | T_i = 0) > 0$ for some (g, u) . Another extreme example is that the total number of treated units is too large. For example, when we use complete randomization with the total number of treated units equal to $N - 1$. In this case, whenever $T_i = 0, \Pr(G = g, U = u | T_i = 0) = 0$ for all (g, u) except for $(g, u) = (1, 1)$, but when $T_i = 1, \Pr(G = g, U = u | T_i = 1) > 0$ for some (g, u) other than $(g, u) = (1, 1)$. It is clear that when researchers use a Bernoulli design, the support of $\Pr(G_i = g, U_i = u | T_i = 1)$ is equal to the support of $\Pr(G_i = g, U_i = u | T_i = 0)$ for all i .

When we define the unit level network-specific spillover effect, we avoid ill-defined causal effects by focusing on settings where $\Pr(U_i = u | T_i = d, G_i = g^H)$ and $\Pr(U_i = u | T_i = d, G_i = g^L)$ have the same support for all i . This requires that g^H and g^L are small enough so that the distribution over the fraction of treated neighbors in network \mathcal{U} is not restricted, especially $\Pr(U_i = 0 | T_i = d, G_i = g^H) > 0$ and $\Pr(U_i = 0 | T_i = d, G_i = g^L) > 0$ for all i . Formally, $g^H, g^L \leq g_s$ where $g_s \equiv \min\{1 - |\mathcal{N}_i^{(\mathcal{G}, \mathcal{U})}| / |\mathcal{N}_i^{\mathcal{G}}|\}$.

The desired support condition can be violated when the total number of treated units is too small so that for some $u, \Pr(U = u | T_i = 1, G_i = g^H) = 0$ and $\Pr(U = u | T_i = 1, G_i = g^L) > 0$. One extreme example is that when we use complete randomization with the total number of treated units equal to $1 + g^H \times |\mathcal{N}_i^{\mathcal{G}}|$. In this case, whenever $G_i = g^H, \Pr(U = u | T_i = 1, G_i = g^H) = 0$ for all $u > |\mathcal{N}_i^{(\mathcal{G}, \mathcal{U})}| / |\mathcal{N}_i^{\mathcal{U}}|$, but when $G_i = g^L < g^H, \Pr(U = u | T_i = 0, G_i = g^L) > 0$ for some $u > |\mathcal{N}_i^{(\mathcal{G}, \mathcal{U})}| / |\mathcal{N}_i^{\mathcal{U}}|$. Finally, it is clear that $\Pr(U_i = u | T_i = d, G_i = g^H)$ and $\Pr(U_i = u | T_i = d, G_i = g^L)$ have the same support for all i if researchers use a Bernoulli design and $g^H, g^L \leq g_s$.

Appendix B. Connection between Total Spillover Effects and Network-Specific Spillover Effects

Here, we connect the ANSE to the popular estimand in the literature. In particular, we show that the ANSE can be seen as the decomposition of the average total spillover effect (Hudgens and Halloran 2008).

First, by extending Hudgens and Halloran (2008) to settings with multiple networks, the individual average potential outcome are defined as follows.

$$\bar{Y}_i(d, g) \equiv \sum_{u \in \Delta_i^u} Y_i(d, g, u) \Pr(U_i = u | T_i = d, G_i = g), \tag{A1}$$

where the potential outcome of individual i is averaged over the conditional distribution of the treatment assignment $\Pr(U_i = u | T_i = d, G_i = g)$. Here, the individual average potential outcome represents the expected outcome of unit i when she receives the direct treatment d and the treated proportion g in network \mathcal{G} . Taking the difference in the two individual average potential outcomes, the *average total spillover effect* (ATSE) in network \mathcal{G} is defined as follows (Halloran and Hudgens 2016).⁸

$$\psi(g^H, g^L; d) \equiv \frac{1}{N} \sum_{i=1}^N \{\bar{Y}_i(d, g^H) - \bar{Y}_i(d, g^L)\}. \tag{A2}$$

This causal quantity is the *total* spillover effect of changing the treated proportion in network \mathcal{G} from g^L to g^H as the following decomposition of the ATSE demonstrates.

$$\psi(g^H, g^L; d) = \tau(g^H, g^L; d) + \tag{A3}$$

⁸ This quantity is called the average indirect causal effect in Hudgens and Halloran (2008). We define it as the average total spillover effect to clarify how it combines multiple network-specific spillover effects.

$$\frac{1}{N} \sum_{i=1}^N \left\{ \sum_{u \in \Delta_i^u} \{Y_i(d, g^H, u) - Y_i(d, g^L, u)\} \{ \Pr(U_i = u \mid T_i = d, G_i = g^H) - \Pr(U_i = u \mid T_i = d, G_i = g^L) \} \right\},$$

for any $u' \in \Delta_i^u$. The first term is the ANSE in network \mathcal{G} (Definition 3), which quantifies the spillover effect specific to network \mathcal{G} . The second term represents the spillover effect in \mathcal{U} , $Y_i(d, g^H, u) - Y_i(d, g^L, u)$, weighted by the change in the conditional distribution of U_i due to the change in G_i , $\Pr(U_i = u \mid T_i = d, G_i = g^H) - \Pr(U_i = u \mid T_i = d, G_i = g^L)$. This is because U_i , the treated proportion of neighbors in the other network \mathcal{U} , is not fixed at constant and thus, they change as G_i , the treated proportion of neighbors in \mathcal{G} , changes. Thus, the ATSE captures the sum of the spillover effect specific to network \mathcal{G} and the spillover effect specific to \mathcal{U} induced by the change in U_i associated with the change in G_i . For example, the ATSE of changing from g^L to g^H on the Facebook network captures two spillover effects together; (1) the spillover effect specific to the Facebook and (2) the spillover effect in the face-to-face network. This is because the treated proportion in the offline network U_i is associated with the change in the treated proportion in the Facebook network G_i . We discuss this issue in further details when we derive the exact bias formula in Section 3. When network \mathcal{U} , such as the offline network, is causally irrelevant, the ATSE is equal to the ANSE in the Facebook network, but in general, the two estimands do not coincide.

While both the ATSE and the ANSE quantify spillover effects, their substantive meanings differ. The ATSE is useful when researchers wish to know the total amount of spillover effects that result from interventions on an observed network. For instance, politicians decided to run online campaigns on Twitter and want to estimate the total amount of spillover effects they can induce by their Twitter messages. These politicians might not be interested in distinguishing whether the spillover effects arise through Twitter or through unobserved face-to-face interactions. Thus, the ATSE is of relevance when the target network is predetermined and the mechanism can be ignored.

In contrast, the ANSE is essential for disentangling different channels through which spillover effects arise. It is the main quantity of interest when researchers wish to examine the causal role of individual networks or to discover the most causally relevant network to target. For example, it is of scientific interest to distinguish how much spillover effects arise through the Twitter network or through offline communications. By estimating the ANSE, researchers can learn about the importance of online human interactions.

Appendix C. Proofs

This section provides proofs for all theorems in the paper.

C.1 Proof of Theorem 1

C.1.1 Average Direct Effect. First, we rewrite the estimator with the standard IPW representation.

$$\begin{aligned} \hat{\delta} &= \frac{1}{N} \sum_{i=1}^N 1\{T_i = 1\} \tilde{w}_i Y_i - \frac{1}{N} \sum_{i=1}^N 1\{T_i = 0\} \tilde{w}_i Y_i \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{(g,u) \in \Delta_i^{gu}} \Pr(G_i = g, U_i = u) \left\{ \frac{1\{T_i = 1, G_i = g, U_i = u\} Y_i}{\Pr(T_i = 1, G_i = g, U_i = u)} - \frac{1\{T_i = 0, G_i = g, U_i = u\} Y_i}{\Pr(T_i = 0, G_i = g, U_i = u)} \right\}. \end{aligned}$$

Then, the theorem follows from the standard proof for the IPW estimator.

$$\begin{aligned}
 & \mathbb{E}[\hat{\delta}] \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{(g,u) \in \Delta_i^{gu}} \Pr(G_i = g, U_i = u) \times \\
 & \quad \left\{ \frac{\mathbb{E}[1\{T_i = 1, G_i = g, U_i = u\}Y_i]}{\Pr(T_i = 1, G_i = g, U_i = u)} - \frac{\mathbb{E}[1\{T_i = 0, G_i = g, U_i = u\}Y_i]}{\Pr(T_i = 0, G_i = g, U_i = u)} \right\} \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{(g,u) \in \Delta_i^{gu}} \Pr(G_i = g, U_i = u) \times \\
 & \quad \left\{ \frac{\Pr(T_i = 1, G_i = g, U_i = u)Y_i(1, g, u)}{\Pr(T_i = 1, G_i = g, U_i = u)} - \frac{\Pr(T_i = 0, G_i = g, U_i = u)Y_i(0, g, u)}{\Pr(T_i = 0, G_i = g, U_i = u)} \right\} \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{(g,u) \in \Delta_i^{gu}} \Pr(G_i = g, U_i = u) \{Y_i(1, g, u) - Y_i(0, g, u)\} \\
 &= \delta
 \end{aligned}$$

where the second equality follows from the consistency of potential outcomes. □

C.1.2 Average Network-Specific Effect. First, we rewrite the estimator with the standard IPW representation.

$$\begin{aligned}
 & \hat{\tau} \\
 &= \frac{1}{N} \sum_{i=1}^N 1\{T_i = d, G_i = g^H\} w_i Y_i - \frac{1}{N} \sum_{i=1}^N 1\{T_i = d, G_i = g^L\} w_i Y_i \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \Pr(U_i = u | T_i = d, G_i = g^L) \times \\
 & \quad \left\{ \frac{1\{T_i = d, G_i = g^H, U_i = u\}Y_i}{\Pr(T_i = d, G_i = g^H, U_i = u)} - \frac{1\{T_i = d, G_i = g^L, U_i = u\}Y_i}{\Pr(T_i = d, G_i = g^L, U_i = u)} \right\},
 \end{aligned}$$

Then, the theorem follows from the standard proof for the IPW estimator.

$$\begin{aligned}
 & \mathbb{E}[\hat{\tau}(g, g'; d)] \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \Pr(U_i = u | T_i = d, G_i = g^L) \times \\
 & \quad \left\{ \frac{\mathbb{E}[1\{T_i = d, G_i = g^H, U_i = u\}Y_i]}{\Pr(T_i = d, G_i = g^H, U_i = u)} - \frac{\mathbb{E}[1\{T_i = d, G_i = g^L, U_i = u\}Y_i]}{\Pr(T_i = d, G_i = g^L, U_i = u)} \right\} \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \Pr(U_i = u | T_i = d, G_i = g^L) \times \\
 & \quad \left\{ \frac{\Pr(T_i = d, G_i = g^H, U_i = u)Y_i(d, g^H, u)}{\Pr(T_i = d, G_i = g^H, U_i = u)} - \frac{\Pr(T_i = d, G_i = g^L, U_i = u)Y_i(d, g^L, u)}{\Pr(T_i = d, G_i = g^L, U_i = u)} \right\} \\
 &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \Pr(U_i = u | T_i = d, G_i = g^L) \{Y_i(d, g^H, u) - Y_i(d, g^L, u)\} \\
 &= \tau(g^H, g^L; d),
 \end{aligned}$$

which completes the proof. □

C.2 Proof of Theorem 2

The expectation of an estimator $\hat{\tau}_B(\mathbf{g}^H, \mathbf{g}^L; d)$ is

$$\begin{aligned} \mathbb{E}[\hat{\tau}_B(\mathbf{g}^H, \mathbf{g}^L; d)] &= \frac{1}{N} \sum_{i=1}^N \left\{ \mathbb{E}[Y_i \mid T_i = d, G_i = \mathbf{g}^H] - \mathbb{E}[Y_i \mid T_i = d, G_i = \mathbf{g}^L] \right\} \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \left\{ Y_i(d, \mathbf{g}^H, u) \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^H) - Y_i(d, \mathbf{g}^L, u) \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^L) \right\}. \end{aligned}$$

Therefore, we get

$$\begin{aligned} &\mathbb{E}[\hat{\tau}_B(\mathbf{g}^H, \mathbf{g}^L; d)] - \tau(\mathbf{g}^H, \mathbf{g}^L; d) \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \left\{ Y_i(d, \mathbf{g}^H, u) \{ \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^H) - \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^L) \} \right. \\ &\quad \left. - Y_i(d, \mathbf{g}^L, u) \{ \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^L) - \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^H) \} \right\} \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{u \in \Delta_i^u} \left\{ \{ Y_i(d, \mathbf{g}^H, u) - Y_i(d, \mathbf{g}^L, u) \} \right. \\ &\quad \left. \times \{ \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^H) - \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^L) \} \right\}. \end{aligned}$$

for any $u' \in \Delta^u$. □

C.2.1 Lemma: Bias in ADE. First, we can rewrite the estimator as the standard IPW estimator.

$$\begin{aligned} \hat{\delta}_B &= \frac{1}{N} \sum_{i=1}^N 1\{T_i = 1\} \tilde{w}_i^B Y_i - \frac{1}{N} \sum_{i=1}^N 1\{T_i = 0\} \tilde{w}_i^B Y_i \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{g \in \Delta_i^g} \Pr(G_i = g) \left\{ \frac{1\{T_i = 1, G_i = g\} Y_i}{\Pr(T_i = 1, G_i = g)} - \frac{1\{T_i = 0, G_i = g\} Y_i}{\Pr(T_i = 0, G_i = g)} \right\}. \end{aligned}$$

We have the following equality for any g ,

$$\begin{aligned} &\mathbb{E}[1\{T_i = d, G_i = g\} Y_i] \\ &= \mathbb{E} \left[\sum_{u \in \Delta_i^u(g)} 1\{T_i = d, G_i = g, U_i = u\} Y_i(d, g, u) \right] \\ &= \sum_{u \in \Delta_i^u(g)} \Pr(T_i = d, G_i = g, U_i = u) Y_i(d, g, u) \end{aligned}$$

where $\Delta_i^u(g)$ is the support $\{u : \Pr(U_i = u \mid G_i = g) > 0\}$. Therefore, the expectation of $\hat{\delta}_B$ is

$$\begin{aligned} &\mathbb{E}[\hat{\delta}_B] \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{g \in \Delta_i^g} \Pr(G_i = g) \left\{ \frac{\mathbb{E}[1\{T_i = 1, G_i = g\} Y_i]}{\Pr(T_i = 1, G_i = g)} - \frac{\mathbb{E}[1\{T_i = 0, G_i = g\} Y_i]}{\Pr(T_i = 0, G_i = g)} \right\} \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{g \in \Delta_i^g} \Pr(G_i = g) \end{aligned}$$

$$\begin{aligned} & \left\{ \sum_{u \in \Delta_i^u(g)} \left\{ \frac{\Pr(T_i = 1, G_i = g, U_i = u) Y_i(1, g, u)}{\Pr(T_i = 1, G_i = g)} - \frac{\Pr(T_i = 0, G_i = g, U_i = u) Y_i(0, g, u)}{\Pr(T_i = 0, G_i = g)} \right\} \right\} \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{g \in \Delta_i^g} \Pr(G_i = g) \\ & \left\{ \sum_{u \in \Delta_i^u(g)} \left\{ Y_i(1, g, u) \Pr(U_i = u | T_i = 1, G_i = g) - Y_i(0, g, u) \Pr(U_i = u | T_i = 0, G_i = g) \right\} \right\}. \end{aligned}$$

Then, we have

$$\begin{aligned} & \mathbb{E}[\hat{\delta}_B] - \delta \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{g \in \Delta_i^g} \Pr_i(G_i = g) \left\{ \sum_{u \in \Delta_i^u} Y_i(1, g, u) \{ \Pr_i(U_i = u | T_i = 1, G_i = g) - \Pr_i(U_i = u | G_i = g) \} \right. \\ & \quad \left. - \sum_{u \in \Delta_i^u} Y_i(0, g', u) \{ \Pr(U_i = u | T_i = 0, G_i = g) - \Pr(U_i = u | G_i = g) \} \right\} \\ &= \frac{1}{N} \sum_{i=1}^N \sum_{g \in \Delta_i^g} \Pr(G_i = g) \\ & \left\{ \sum_{u \in \Delta_i^u(g)} \{ Y_i(1, g, u) - Y_i(1, g, u') \} \{ \Pr(U_i = u | T_i = 1, G_i = g) - \Pr(U_i = u | G_i = g) \} \right. \\ & \quad \left. - \sum_{u \in \Delta_i^u(g)} \{ Y_i(0, g', u) - Y_i(0, g', u') \} \{ \Pr(U_i = u | T_i = 0, G_i = g) - \Pr(U_i = u | G_i = g) \} \right\}, \end{aligned}$$

which completes the proof. □

C.3 Proof of Theorem 3

Using Theorem 2, under Assumption 3,

$$\begin{aligned} & \mathbb{E}[\hat{\tau}_B(g^H, g^L; d)] - \tau(g^H, g^L; d) \\ &= \lambda \times \frac{1}{N} \sum_{i=1}^N \{ \mathbb{E}[U_i = u | T_i = d, G_i = g^H] - \mathbb{E}[U_i = u | T_i = d, G_i = g^L] \}. \end{aligned}$$

From here, we focus on $\mathbb{E}[U_i | T_i = d, G_i = g^H]$. For notational simplicity, we use $n_G(i)$ to denote the number of neighbors in the network \mathcal{G} for individual i and $n_U(i)$ is similarly defined. Also, for individual i , let $\pi_{GU}(i)$ be the fraction of the neighbors in \mathcal{U} who are neighbors in \mathcal{G} as well. Formally, $n_G(i) = |\mathcal{N}_i^{\mathcal{G}}|$, $n_U(i) = |\mathcal{N}_i^{\mathcal{U}}|$ and $\pi_{GU}(i) = |\mathcal{N}_i^{(\mathcal{G}, \mathcal{U})}| / |\mathcal{N}_i^{\mathcal{U}}|$.

First, we consider Bernoulli randomization with probability p . Under this setting,

$$\mathbb{E}[U_i | T_i = t, G_i = g^H] = \pi_{GU}(i) \times g^H + (1 - \pi_{GU}(i)) \times p.$$

Therefore, we have

$$\mathbb{E}[\hat{\tau}_B(g^H, g^L; d)] - \tau(g^H, g^L; d)$$

$$\begin{aligned}
 &= \lambda \times \frac{1}{N} \sum_{i=1}^N \{ \mathbb{E}[U_i = u \mid T_i = d, G_i = g^H] - \mathbb{E}[U_i = u \mid T_i = d, G_i = g^L] \}. \\
 &= \lambda \times \frac{1}{N} \sum_{i=1}^N \pi_{GU}(i) \times (g^H - g^L) \\
 &= \lambda \times \pi_{GU} \times (g^H - g^L).
 \end{aligned}$$

where the final equality follows from the definition of π_{GU} .

Next, we consider complete randomization with the number of treated units K . Under this setting,

$$\begin{aligned}
 &\mathbb{E}[U_i \mid T_i = d, G_i = g^H] \\
 &= \frac{n_U(i) \times \pi_{GU}(i) \times g^H}{n_U(i)} + (1 - \pi_{GU}(i)) \frac{K - d - n_G(i) \times g^H}{N - 1 - n_G(i)} \\
 &= \pi_{GU}(i) \times g^H + (1 - \pi_{GU}(i)) \frac{K - d - n_G(i) \times g^H}{N - 1 - n_G(i)} \\
 &= \left\{ \pi_{GU}(i) - \frac{n_G(i)}{N - 1 - n_G(i)} (1 - \pi_{GU}(i)) \right\} g^H + \frac{K - d}{N - 1 - n_G(i)} (1 - \pi_{GU}(i))
 \end{aligned}$$

When N is much larger than $n_G(i)$, $n_G(i)/(N - 1 - n_G(i)) \approx 0$. Then, we have

$$\begin{aligned}
 \mathbb{E}[U_i \mid T_i = d, G_i = g^H] &\approx \pi_{GU}(i) g^H + \frac{K - d}{N - 1 - n_G(i)} (1 - \pi_{GU}(i)), \\
 \mathbb{E}[U_i \mid T_i = d, G_i = g^H] - \mathbb{E}[U_i = u \mid T_i = d, G_i = g^L] &\approx \pi_{GU}(i) (g^H - g^L).
 \end{aligned}$$

Therefore, when N is much larger than $n_G(i)$ for all i , we get the simplified bias formula.

$$\begin{aligned}
 &\mathbb{E}[\hat{\tau}_B(g^H, g^L; d)] - \tau(g^H, g^L; d) \\
 &= \lambda \times \frac{1}{N} \sum_{i=1}^N \{ \mathbb{E}[U_i = u \mid T_i = d, G_i = g^H] - \mathbb{E}[U_i = u \mid T_i = d, G_i = g^L] \}. \\
 &\approx \lambda \times \frac{1}{N} \sum_{i=1}^N \pi_{GU}(i) \times (g^H - g^L) \\
 &= \lambda \times \pi_{GU} \times (g^H - g^L).
 \end{aligned}$$

Finally, we consider a situation when N is not large enough to have the aforementioned approximation. Suppose $N \approx (C + 1)n_G(i) + 1$ for all i . Then,

$$\begin{aligned}
 \mathbb{E}[U_i \mid T_i = d, G_i = g^H] &\approx \left\{ \frac{C + 1}{C} \pi_{GU}(i) - \frac{1}{C} \right\} \times g^H + \frac{K - d}{N - 1 - n_G(i)} (1 - \pi_{GU}(i)), \\
 \mathbb{E}[U_i \mid T_i = d, G_i = g^H] - \mathbb{E}[U_i = u \mid T_i = d, G_i = g^L] &\approx \left\{ \frac{C + 1}{C} \pi_{GU}(i) - \frac{1}{C} \right\} (g^H - g^L).
 \end{aligned}$$

Therefore, the bias can be written as,

$$\begin{aligned}
 &\mathbb{E}[\hat{\tau}_B(g^H, g^L; d)] - \tau(g^H, g^L; d) \\
 &= \lambda \times \frac{1}{N} \sum_{i=1}^N \{ \mathbb{E}[U_i = u \mid T_i = d, G_i = g^H] - \mathbb{E}[U_i = u \mid T_i = d, G_i = g^L] \}.
 \end{aligned}$$

$$\begin{aligned} &\approx \lambda \times \left\{ \frac{C+1}{C} \times \frac{1}{N} \sum_{i=1}^N \pi_{GU}(i) - \frac{1}{C} \right\} \times (g^H - g^L). \\ &= \lambda \times \left\{ \frac{C+1}{C} \pi_{GU} - \frac{1}{C} \right\} \times (g^H - g^L). \end{aligned} \tag{A4}$$

□

C.4 Proof of Theorem 4

First, we set the following notations. We define the support Δ_s^u to be the support Δ_i^u for all i with $S_i = s$. We drop subscript s whenever it is obvious from contexts. For $\bar{g} \in \{g^H, g^L\}$,

$$\begin{aligned} r_{\bar{g}}(u) &\equiv \frac{1}{N} \sum_{i: S_i=s} Y_i(d, \bar{g}, u) \\ v_{g^H}(\bar{g}) &\equiv \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\max_u r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)} \\ v_{g^L}(\bar{g}) &\equiv \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\max_u r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)} \\ \Gamma(\bar{g}) &\equiv \frac{v_{g^H}(\bar{g})}{v_{g^L}(\bar{g})} = \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L)} \\ MR^{obs}(g^H, g^L; s) &\equiv \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)} \\ MR_{\bar{g}}^{true}(g^H, g^L; s) &\equiv \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = \bar{g})}{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = \bar{g})} \end{aligned}$$

where $0 \leq v_{g^H}(\bar{g}), v_{g^L}(\bar{g}) \leq 1$ because of non-negative outcomes.

Lemma 1 For (g^H, g^L) ,

$$\begin{aligned} \frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)} &\leq B \quad \frac{MR^{obs}(g^H, g^L; s)}{MR_{g^H}^{true}(g^H, g^L; s)} \leq B, \\ \frac{MR^{obs}(g^L, g^H; s)}{MR_{g^L}^{true}(g^L, g^H; s)} &\leq B \quad \frac{MR^{obs}(g^L, g^H; s)}{MR_{g^H}^{true}(g^L, g^H; s)} \leq B. \end{aligned}$$

Proof. This proof closely follows Ding and VanderWeele (2016). The key difference is that we study bias due to an unmeasured relevant network in the presence of interference in multiple networks in contrary to bias due to an unmeasured confounder in observational studies without interference (Ding and VanderWeele 2016).

For $\bar{g} \in \{g^H, g^L\}$ and s ,

$$\begin{aligned} \Gamma(\bar{g}) &= \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L)} \\ &= \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \frac{\Pr(U_i = u \mid T_i = d, G_i = g^H)}{\Pr(U_i = u \mid T_i = d, G_i = g^L)} \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L)} \\ &\leq RR_{GU} \end{aligned}$$

Also, for $\bar{g} \in \{g^H, g^L\}$ and s ,

$$\begin{aligned} \Gamma(\bar{g}) &= \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^H)} \\ &= \frac{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \frac{\Pr(U_i = u \mid T_i = d, G_i = g^L)}{\Pr(U_i = u \mid T_i = d, G_i = g^H)} \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} \{r_{\bar{g}}(u) - \min_u r_{\bar{g}}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^H)} \\ &\leq RR_{GU}. \end{aligned}$$

Then, we have

$$\begin{aligned} &\frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)} \\ &= \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)} \times \frac{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)} \\ &= \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)} \\ &= \frac{\{\max_u r_{g^H}(u) - \min_u r_{g^H}(u)\} v_{g^H}(g^H) + \min_u r_{g^H}(u)}{\{\max_u r_{g^H}(u) - \min_u r_{g^H}(u)\} \frac{v_{g^H}(g^H)}{\Gamma(g^H)} + \min_u r_{g^H}(u)} \end{aligned}$$

From Lemma A.1 in Ding and VanderWeele (2016), when $\Gamma(g^H) > 1$, $\frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)}$ is increasing in $v_{g^H}(g^H)$. Therefore, it takes the maximum value when $v_{g^H}(g^H) = 1$.

$$\begin{aligned} \frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)} &\leq \frac{\Gamma(g^H) \times MR_{UY}(g^H, s)}{\Gamma(g^H) + MR_{UY}(g^H, s) - 1} \\ &\leq \frac{RR_{GU} \times MR_{UY}}{RR_{GU} + MR_{UY} - 1} \end{aligned}$$

where the second inequality comes from Lemma A.2 in Ding and VanderWeele (2016) and $\Gamma(g^H) \leq RR_{GU}$, $MR_{UY} = \max_{g,s} MR_{UY}(g, s)$.

From Lemma A.1 in Ding and VanderWeele (2016), when $\Gamma(g^H) \leq 1$, $\frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)}$ is nonincreasing in $v_{g^H}(g^H)$. Therefore, it takes the maximum value at $v_{g^H}(g^H) = 0$.

$$\frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)} \leq 1 \leq \frac{RR_{GU} \times MR_{UY}}{RR_{GU} + MR_{UY} - 1}$$

where the second inequality comes from Lemma A.2 in Ding and VanderWeele (2016) and $RR_{GU} \geq 1$, $MR_{UY} \geq 1$.

Hence, we obtain the desired result.

$$\frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)} \leq \frac{RR_{GU} \times MR_{UY}}{RR_{GU} + MR_{UY} - 1}.$$

Similar derivations apply to the other three inequalities. □

Proof of the theorem. For notational simplicity, we use the following representation.

$$\begin{aligned}
 m(d, g; s) &\equiv \frac{1}{N} \sum_{i: S_i = s} \mathbb{E}[Y_i | T_i = d, G_i = g] \\
 &= \frac{1}{N} \sum_{i: S_i = s} \frac{\sum_{u \in \Delta_s^u} \Pr(T_i = d, G_i = g, U_i = u) Y_i(d, g, u)}{\Pr(T_i = d, G_i = g)} \\
 &= \frac{1}{N} \sum_{i: S_i = s} \sum_{u \in \Delta_s^u} \Pr(U_i = u | T_i = d, G_i = g) Y_i(d, g, u) \\
 &= \sum_{u \in \Delta_s^u} \left\{ \frac{1}{N} \sum_{i: S_i = s} Y_i(d, g, u) \right\} \Pr(U_i = u | T_i = d, G_i = g) \\
 &= \sum_{u \in \Delta_s^u} r_g(u) \Pr(U_i = u | T_i = d, G_i = g).
 \end{aligned}$$

We want to show that, for g^H, g^L ,

$$\frac{m(d, g^H; s)}{B} - B \times m(d, g^L; s) \leq \frac{1}{N} \sum_{i: S_i = s} \tau_i(g^H, g^L; d) \leq B \times m(d, g^H; s) - \frac{m(d, g^L; s)}{B}.$$

Because this implies the desired result.

$$\begin{aligned}
 \frac{m(d, g^H; s)}{B} - B \times m(d, g^L; s) &\leq \frac{1}{N} \sum_{i: S_i = s} \tau_i(g^H, g^L; d) \leq B \times m(d, g^H; s) - \frac{m(d, g^L; s)}{B} \\
 \Leftrightarrow \left\{ \begin{array}{l} \sum_{s \in \mathcal{S}} \left\{ \frac{m(d, g^H; s)}{B} - B \times m(d, g^L; s) \right\} \leq \sum_{s \in \mathcal{S}} \left\{ \frac{1}{N} \sum_{i: S_i = s} \tau_i(g^H, g^L; d) \right\} \\ \sum_{s \in \mathcal{S}} \left\{ \frac{1}{N} \sum_{i: S_i = s} \tau_i(g^H, g^L; d) \right\} \leq \sum_{s \in \mathcal{S}} \left\{ B \times m(d, g^H; s) - \frac{m(d, g^L; s)}{B} \right\} \end{array} \right\} \\
 \Leftrightarrow \frac{\mathbb{E}[\hat{m}(d, g^H)]}{B} - B \times \mathbb{E}[\hat{m}(d, g^L)] &\leq \tau(g^H, g^L; d) \leq B \times \mathbb{E}[\hat{m}(d, g^H)] - \frac{\mathbb{E}[\hat{m}(d, g^L)]}{B}.
 \end{aligned}$$

First, using Lemma 1,

$$\begin{aligned}
 &\frac{m(d, g^H; s)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u | T_i = d, G_i = g^L)} \\
 &= \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u | T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u | T_i = d, G_i = g^L)} \\
 &= \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u | T_i = d, G_i = g^H)}{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u | T_i = d, G_i = g^L)} \times \frac{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u | T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u | T_i = d, G_i = g^L)} \\
 &= \frac{MR^{obs}(g^H, g^L; s)}{MR_{g^L}^{true}(g^H, g^L; s)} \leq B
 \end{aligned}$$

where the final equality follows from the lemma. Therefore,

$$\frac{m(d, g^H; s)}{B} \leq \sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u | T_i = d, G_i = g^L). \tag{A5}$$

Also, since $B \geq 1$,

$$\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u | T_i = d, G_i = g^L) = m(d, g^L; s) \leq B \times m(d, g^L; s). \tag{A6}$$

Finally, taking Equations (A5) and (A6) together,

$$\begin{aligned}
 & \frac{m(d, g^H; s)}{B} - B \times m(d, g^L; s) \\
 & \leq \sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L) - \sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L) \\
 & = \sum_{u \in \Delta_s^u} \{r_{g^H}(u) - r_{g^L}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L) \\
 & = \frac{1}{N} \sum_{i: S_i = s} \sum_{u \in \Delta_s^u} \{Y_i(d, g^H, u) - Y_i(d, g^L, u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L) \\
 & = \frac{1}{N} \sum_{i: S_i = s} \tau_i(g^H, g^L; d).
 \end{aligned}$$

Similarly, we want to prove

$$\frac{1}{N} \sum_{i: S_i = s} \tau_i(g^H, g^L; d) \leq B \times m(d, g^H; s) - \frac{m(d, g^L; s)}{B}.$$

First, since $B \geq 1$,

$$\frac{m(d, g^L; s)}{B} \leq m(d, g^L; s) = \sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L). \tag{A7}$$

Then, using Lemma 1,

$$\begin{aligned}
 & \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)}{m(d, g^H; s)} \\
 & = \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^H)} \\
 & = \frac{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)} \times \frac{\sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L)}{\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^H)} \\
 & = \frac{MR^{obs}(g^L, g^H; s)}{MR_{g^L}^{true}(g^L, g^H; s)} \leq B.
 \end{aligned}$$

Therefore, we have

$$\sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L) \leq B \times m(d, g^H; s). \tag{A8}$$

Finally, taking Equations (A7) and (A8) together,

$$\begin{aligned}
 & B \times m(d, g^H; s) - \frac{m(d, g^L; s)}{B} \\
 & \geq \sum_{u \in \Delta_s^u} r_{g^H}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L) - \sum_{u \in \Delta_s^u} r_{g^L}(u) \Pr(U_i = u \mid T_i = d, G_i = g^L) \\
 & = \sum_{u \in \Delta_s^u} \{r_{g^H}(u) - r_{g^L}(u)\} \Pr(U_i = u \mid T_i = d, G_i = g^L)
 \end{aligned}$$

$$\begin{aligned}
&= \frac{1}{N} \sum_{i:S_i=s} \sum_{u \in \Delta_s^u} \{Y_i(d, \mathbf{g}^H, u) - Y_i(d, \mathbf{g}^L, u)\} \Pr(U_i = u \mid T_i = d, G_i = \mathbf{g}^L) \\
&= \frac{1}{N} \sum_{i:S_i=s} \tau_i(\mathbf{g}^H, \mathbf{g}^L; d).
\end{aligned}$$

Hence we have

$$\frac{m(d, \mathbf{g}^H; s)}{B} - B \times m(d, \mathbf{g}^L; s) \leq \frac{1}{N} \sum_{i:S_i=s} \tau_i(\mathbf{g}^H, \mathbf{g}^L; d) \leq B \times m(d, \mathbf{g}^H; s) - \frac{m(d, \mathbf{g}^L; s)}{B},$$

which completes the proof. \square

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Data Availability Statement

The replication materials are available as Egami (2020).

Supplementary Materials

To view supplementary material for this article, please visit <http://dx.doi.org/10.1017/pan.2020.28>.

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