

Causal Mediation Analysis for Difference-in-Difference Design and Panel Data

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Abstract

Advantages of panel data, i.e., difference in difference (DID) design data, are a large sample size and easy availability. Therefore, panel data are widely used in epidemiology and in all social science fields. The literatures on causal inferences of panel data setting or DID design are growing, but no theory or mediation analysis method has been proposed for such settings. In this study, we propose a methodology for conducting causal mediation analysis in DID design and panel data setting. We provide formal counterfactual definitions for controlled direct effect and natural direct and indirect effect in panel data setting and DID design, including the identification and required assumptions. We also demonstrate that, under the assumptions of linearity and additivity, controlled direct effects can be estimated by contrasting marginal and conditional DID estimators whereas natural indirect effects can be estimated by calculating the product of the exposure-mediator DID estimator and the mediator-outcome DID estimator. A panel regression-based approach is also proposed. The proposed method is then used to investigate mechanisms of the effects of the Covid 19 pandemic on the mental health

status of the population. The results revealed that mobility restrictions mediated approximately 45 % of the causal effect of Covid 19 on mental health status.

Introduction

Panel data is a form of dataset widely used in all social science fields, including sociology [1], economics [2, 3], and political science [4-7]. In epidemiology [8-11], panel data setting is usually treated as a quasi-experimental study to evaluate the role of a policy implementation in a specific health-related outcome. If only considering two time points (pretreatment and posttreatment time-points), panel data setting is also known as “difference-in-difference (DID) design” [10]. The panel data format (or DID design format) is the most common format for open access data, which usually includes nationwide or even worldwide information along with repeated observations between groups. For example, the scientific question of interest in this study was the causal effect of daily COVID-19 deaths on mental health status in a national population and the mediating effect (also termed as indirect effect) mediated through mobility restriction. The dataset is in the form of panel data containing information about daily COVID-19 deaths in 40 countries in the past 1 year. The literature on causal inferences of panel data settings or DID design have grown in the past decade. However, all methods proposed so far have only been used to investigate the average causal effect of a treatment among the treated (ATT) instead of the average causal effect in a national population, which is of interest and importance in epidemiology. In addition, methods for investigating causal mechanism under this setting are also required. Causal mediation analysis is a popular technique for investigating causal pathways through mediators of interest [12, 13]. Various studies have proposed methodologies for conducting mediation analysis under different settings [14-18]. However, no method of performing causal mediation analysis of panel data has been reported.

Therefore, this study developed a method of causal mediation analysis in which the causal effect of the overall population is decomposed into two parts: a mediating part and a non-mediating part. We provide formal definitions for causal effects of interest based on counterfactual models. We then provide the required assumptions for identification and estimation. We show that all causal effects can be expressed in terms of traditional DID estimators. Additionally, we illustrate that, when linearity is assumed, the two widely used methods, i.e., difference method and product method, are valid and correspond to different causal interpretations [19].

Our method is motivated by research on the effect of the nationwide COVID-19 outbreak on mental health and its mechanism. Previous studies have investigated how the COVID-19 outbreak affects mental health in the general population [20, 21]. We were interested in the extent to which this effect is mediated by mobility restriction in the general population. Pandemic-induced restrictions on mobility might result from fear of infection or from

government policy. Some researchers have hypothesized that restrictions on various activities and constant inconvenience in daily life likely result in feelings of loneliness and social isolation, which can then have negative mental health impacts, e.g., insomnia [22]. As a result, we hypothesized that restricted mobility is an important mediator of the causal effect of the national COVID-19 outbreak on mental health status. We obtained panel data for the extent of the national COVID-19 outbreak, the extent of mobility restriction, and the mental health status of the population. The Illustration section then demonstrates an application of causal mediation analysis.

Methods

Notations and causal structures

The variables from the panel data structure are labeled with time and group. For a unit g ($g=1, 2, \dots, G$), let A_g denote the exposure of interest. At time t , let M_{gt} denote a potential mediator, and let Y_{gt} denote the outcome. In time t , M_{gt} is assumed to occur earlier than Y_{gt} . Moreover, Y_{g0} and M_{g0} denote baseline values (i.e., “pretreatment values” in previous works) of Y_{gt} and M_{gt} , respectively. Here we only consider two time points: time t (“post-treatment period” in previous works) and the pre-treatment period. In our study, A_g is the extent of the national COVID-19 outbreak, M_{gt} is the extent of mobility restriction, and Y_{gt} is the mental health status of the general population. In this example, the question is how to quantify the causal effect of the extent of the national COVID-19 outbreak A_g on mental health status Y_{gt} in the general population and the extent to which this effect is mediated by mobility restriction. For simplicity, we dichotomize A_g and M_{gt} as follows: $A_g = 0$ denotes that the number of cases of COVID-19 in country g is relatively low compared to the median number of deaths in country g over time; $A_g = 1$ denotes that the extent of the national COVID-19 outbreak is relatively high; $M_{gt} = 0$ denotes that mobility restriction is low compared to the median value for mobility restriction in country g over time; $M_{gt} = 1$ denotes that the mobility restrictions are relatively high.

We then use the counterfactual model to define all causal effects of interest [23, 24]. Let $Y_{gt}(a)$ be the hypothetical value of the outcome that would have occurred if A_g had been set to a . Let $Y_{gt}(a, m)$ be the hypothetical value of the outcome that would have occurred if A_g had been set to a and M_{gt} had been set to m . Similarly, let $M_{gt}(a)$ be the hypothetical value of the mediator if A_g had been set to a . Consistency and composition are assumed [25-27] as follows: the value of $Y_{gt}(a)$ is the exact value of Y_{gt} when $A_g = a$; the value of $Y_{gt}(a, m)$ is the observed value of Y_{gt} when $A_g = a$ and $M_{gt} = m$; the value of $M_{gt}(a)$ is

the observed value of M_{gt} when $A_g = a$. Figure 1(a) depicts the causal relations among variables A_g , M_{gt} , and Y_{gt} . A further assumption is that both exposure and mediator had no effect on the variables in pre-treatment period (NEPT) [28], which can be mathematically expressed as $Y_{g0} = Y_{g0}(a)$, $Y_{g0} = Y_{g0}(a, m)$ and $M_{g0} = M_{g0}(A_g = a)$ for each a and m .

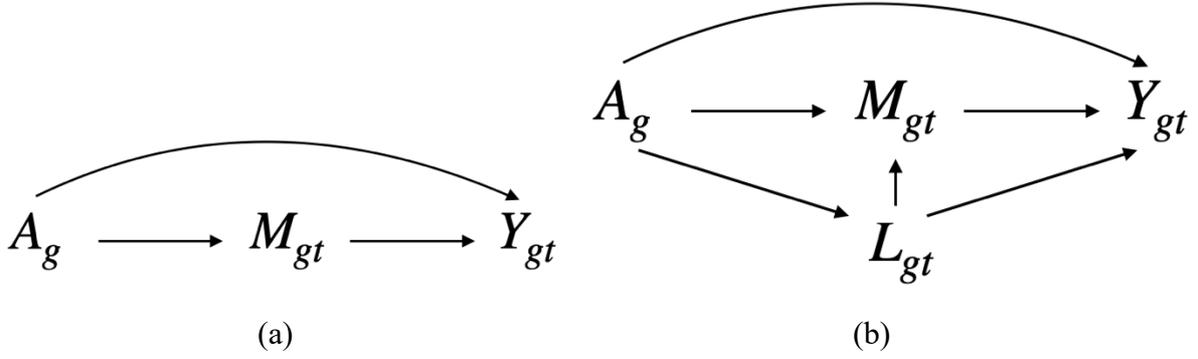


Fig. 1. (a) Causal Diagram of exposure (A_g), mediator (M_{gt}), and outcome (Y_{gt}). (b) Causal Diagram in the presence of “recanting witness” (i.e., exposure-inducing-mediator-outcome confounder) (L_{gt})

General common trend assumption, unconfoundedness and exchangeability in DID design and panel data

Before introducing the causal effects of interest, we discuss the common trend assumption, which is required for identification [9, 10, 29]. The common trend assumption is that, in a counterfactual world in which no one has been exposed, changes in outcome over time are same for all exposure levels. For example, the common trend assumption of the effect from A_g to Y_{gt} implies that $E[Y_{gt}(0) - Y_{g0}(0)|A_g = 0] = E[Y_{gt}(0) - Y_{g0}(0)|A_g = 1]$. Previous literature reveal that, under the common trend assumption, the causal effect of the treatment group (ATT) can be identified [28]. In this study, we are interested in all populations. Therefore, we must apply the common trend assumption in another counterfactual world in which everyone has been exposed, i.e., $E[Y_{gt}(1) - Y_{g0}(1)|A_{gt} = 0] = E[Y_{gt}(1) - Y_{g0}(1)|A_g = 1]$.

A more general assumption that satisfies the above two common trend assumptions is $Y_{gt}(a) - Y_{g0}(a) \perp A_g \forall a$ (CT-0 assumption). That is, the counterfactual trend of Y_{gt} is exchangeable among groups with different values for A_g . This general common trend assumption mathematically equals to $Y_{gt}(a) \perp A_g | Y_{g0} \forall a$ (see Appendix A for detailed proof). This equation also reveals how the common trend assumption is related to the unconfoundedness assumption: the general common trend assumption can be interpreted as the pretreatment outcome (i.e., Y_{g0}) that captures all potential confounders between A_g and Y_{gt} .

Based on the general common trend assumption and its interpretation, we introduce the

assumptions for mediation analysis. We first determine the general common trend assumption of the effect from A_g and M_{gt} to Y_{gt} : (CT-1 assumption) $Y_{gt}(a, m) - Y_{g0}(a, m) \perp A_g \forall a$ and (CT-2 assumption) $Y_{gt}(a, m) - Y_{g0}(a, m) \perp M_{gt} | A_g \forall a, m$. We also determine the general common trend assumption of the effect from A_g to M_{gt} : (CT-3 assumption) $M_{gt}(a) - M_{g0}(a) \perp A_g \forall a$. Similar interpretation of CT-0 can be implied for the above three assumptions (i.e. CT-1, CT-2, and CT-3). The CT-1 assumption and the CT-0 assumption have identical interpretations; the CT-2 assumption can be interpreted as the assumption that both pretreatment outcome and exposure capture all potential confounders between M_{gt} and Y_{gt} ; the CT-3 assumption can be interpreted as the assumption that the pretreatment mediator captures all potential confounders between M_{gt} and A_g .

We also require one additional “cross-world” common trend assumption: (CT-4 assumption) $Y_{gt}(a, m) - Y_{g0}(a, m) \perp M_{gt}(a^*) | A_g \forall a, a^*, m$. The sufficient conditions for this assumption are CT-2 assumption and no mediator-outcome confounder affected by the exposure. This assumption is required for identifying the NDE and NIE even in the non-DID design setting. The exposure-inducing-mediator-outcome confounder is also termed the “recanting witness” in some literature. Figure 1(b) is an example of a CT-4 violation in the presence of a recanting witness (L_{gt}) [30].

The following sections discuss the conventional definitions of total effect, direct effect, and indirect effect in the DID setting and panel data [31]. We also show that, with binary A_g and M_{gt} , all effects can be expressed in terms of traditional DID estimates. It merits noting that the assumptions of consistency, composition, and NEPT are always presumed to hold true. However, the rest of this article only discusses when CT assumptions should be applied.

Definition, identification and estimation of total effect

The total effect (TE) from A_g to Y_{gt} is defined as $E[Y_{gt}(1) - Y_{gt}(0)]$, where the average outcome obtained if A_g had been set to 1 is compared with the average outcome obtained if A_g had been 0. In this study, TE is the effect of the extent of the national COVID-19 outbreak on the mental health status of the general population. Under CT-0 assumption, TE can be identified as DID_{Y_{gt}, A_g} , where $DID_{Y_{gt}, A_g} := \{E[Y_{gt} | A_g = 1] - E[Y_{g0} | A_g = 1]\} - \{E[Y_{gt} | A_g = 0] - E[Y_{g0} | A_g = 0]\}$.

Details of the identification process appear in Appendix B. The DID_{Y_{gt}, A_g} is a DID estimator [11, 32-34]. Notably, previous literature only apply the conventional common trend

assumption and interpret DID_{Y_{gt}, A_g} as the ATT. In contrast, our study applies a general common trend assumption (which is stronger than the traditional one): the identical DID_{Y_{gt}, A_g} can be interpreted as the causal effect for the total population. The DID_{Y_{gt}, A_g} can be used to estimate TE as the difference in the conditional means of Y_{gt} and Y_{g0} where $A_g = 1$ minus that of where $A_g = 0$.

For random variables W and X (where X is binary), a marginal DID estimator can be denoted in simplified notation as $DID_{W,X} := \{E[W_{gt}|X = 1] - E[W_{g0}|X = 1]\} - \{E[W_{gt}|X = 0] - E[W_{g0}|X = 0]\}$. Additionally, for random variables W and X and an additional random variable Z , a conditional DID estimator can be denoted as $DID_{W,X|Z} := \{E[W_{gt}|X = 1, Z] - E[W_{g0}|X = 1, Z]\} - \{E[W_{gt}|X = 0, Z] - E[W_{g0}|X = 0, Z]\}$.

Definition, Identification and Estimation based on DID estimate of Controlled Direct Effect (CDE)

According to different research questions, two decomposition strategies for mediation analysis are available [35]. To evaluate the extent of the effect of an exposure, which is eliminated by the intervening the mediator, TE is decomposed into CDE and portion eliminated (PE) [31, 36], i.e., $TE = CDE + PE$. the extent of effect of the exposure, which can be explained by the mediation mechanisms via the mediator [36, 37], TE is decomposed into natural direct effect (NDE) and natural indirect effect (NIE), i.e. $TE = NDE + NIE$. The causal mediation analysis method proposed in this method is applicable for DID design and panel data setting. This section discusses CDE and PE, and the next section discusses NDE and NIE.

The CDE, which is defined as $CDE(m) = E[Y_{gt}(1, m)] - E[Y_{gt}(0, m)]$, compares exposure between level 1 and level 0 and sets the mediator level to m . The CDE captures the effect of exposure A_g on outcome Y_{gt} by setting M_{gt} to m . Note that CDE may vary with m if effects of A_g interact with effects of M_{gt} . In our motivating example, CDE (1) is interpreted as the change in mental health status caused by the increased severity of the national COVID-19 outbreak when the mobility is always highly restricted by government policy (such as lockdown policy).

Under assumptions CT-1 and CT-2, CDE can be identified as a conditional DID estimator of Y_{gt} and A_g conditioning on M_{gt} as m , $DID_{Y_{gt}, A_g | M_{gt}=m}$, where

$$DID_{Y_{gt}, A_g | M_{gt}=m} := \left[E[Y_{gt} | A_g = 1, M_{gt} = m] - E[Y_{g0} | A_g = 1, M_{gt} = m] \right] - \left[E[Y_{gt} | A_g = 0, M_{gt} = m] - E[Y_{g0} | A_g = 0, M_{gt} = m] \right].$$

Appendix C provides the details of the identification process. The CDE is a function of m , which is set according to the research question. The $\mathbf{DID}_{Y_{gt}, A_g | M_{gt}=m}$ can be used to estimate CDE by calculating the difference in the conditional means of Y_{gt} and Y_{g0} where $A_g = 1$ and $M_{gt} = m$ minus the difference in the conditional means of Y_{gt} and Y_{g0} where $A_g = 0$ and $M_{gt} = m$.

The PE can then be calculated as the difference between TE and CDE as

$$\text{PE}(m) = \text{TE} - \text{CDE}(m) = \mathbf{DID}_{Y_{gt}, A_g} - \mathbf{DID}_{Y_{gt}, A_g | M_{gt}=m},$$

which is the contrast of marginal DID mediator of Y_{gt} and A_g and conditional DID mediator conditional on M_{gt} as m . After TE and CDE are estimated, PE can be estimated.

Definition, Identification and Estimation based on DID estimate of NDE and NIE

Based on the traditional definition of NDE [31, 38], this study defined NDE for DID design and panel data settings as

$$E \left[Y_{gt} \left(1, M_{gt}(0) \right) - Y_{gt} \left(0, M_{gt}(0) \right) \right],$$

the contrast of the expectation of hypothetical values of Y_{gt} had the exposure been set to 1 (versus 0) and had the mediator been always set to the hypothetical value had the exposure been set to 0. In our motivating example, this would capture the change in mental health status between two levels of severity of the national COVID-19 outbreak if the maximum restriction on mobility for each country had always been set to the mobility level in the absence of a national COVID-19 outbreak. The NIE can be defined as $E \left[Y_{gt} \left(1, M_{gt}(1) \right) - Y_{gt} \left(1, M_{gt}(0) \right) \right]$, which is the difference in the hypothetical values of Y_{gt} had the exposure always been set to 1, but had the mediator been set to two hypothetical values had the exposure been set to 1 versus 0. In our example, NIE captures mental health status by comparing the mobility restriction policy under two levels of severity of the national COVID-19 outbreak if the severity of the national COVID-19 outbreak had in fact been 1.

Four common trend assumptions are needed to identify NDE and NIE. Under assumptions CT-1 to 4, NIE can be identified as $\text{NIE} = \mathbf{DID}_{M_{gt}|A_g} \times \mathbf{DID}_{Y_{gt}, M_{gt}|A_g=1}$, where

$$\mathbf{DID}_{M_{gt}, A_g} := \{E[M_{gt}|A_g = 1] - E[M_{g0}|A_g = 1]\} - \{E[M_{gt}|A_g = 0] - E[M_{g0}|A_g = 0]\},$$

and

$$\mathbf{DID}_{Y_{gt}, M_{gt}|A_g=1} := \left[E[Y_{gt}|A_g = 1, M_{gt} = 1] - E[Y_{g0}|A_g = 1, M_{gt} = 1] \right] -$$

$$\left[E[Y_{gt}|A_g = 1, M_{gt} = 0] - E[Y_{g0}|A_g = 1, M_{gt} = 0] \right] \Big\}.$$

The details of the procedures appear in Appendix D. The first term ($\mathbf{DID}_{M_{gt}, A_g}$) is the marginal DID estimator of M_{gt} and A_g , which can be interpreted as the effect of A_g on M_{gt} . The second term ($\mathbf{DID}_{Y_{gt}, M_{gt}|A_g=1}$) is the conditional DID estimator of Y_{gt} and M_{gt} when $A_g = 1$, which can be interpreted as the effect of M_{gt} on Y_{gt} . Therefore, the NIE can be expressed as the product of the effect of A_g on M_{gt} and the effect of M_{gt} on Y_{gt} , expressed in the form of DID estimators. The $\mathbf{DID}_{M_{gt}, A_g}$ can be used to estimate the effect of A_g on M_{gt} based on the difference between the conditional means of M_{gt} and M_{g0} where $A_g = 1$ and the conditional means of M_{gt} and M_{g0} , where $A_g = 0$. Meanwhile, $\mathbf{DID}_{Y_{gt}, M_{gt}|A_g=1}$ can be used to estimate the effect of M_{gt} on Y_{gt} by calculating the difference in the conditional means of Y_{gt} and Y_{g0} where $A_g = 1$ and $M_{gt} = 1$ minus the conditional means of Y_{gt} and Y_{g0} where $A_g = 1$ and $M_{gt} = 0$. The product of estimators $\mathbf{DID}_{M_{gt}, A_g}$ and $\mathbf{DID}_{Y_{gt}, M_{gt}|A_g=1}$ obtains the estimation of NIE. Thus, NDE is the difference between TE and NIE, i.e., $NDE = TE - NIE$. Estimation of TE and NIE enables estimation of NDE.

Empirical 2FE regression-based method

In the above discussion, the mediator was binary, and effects were only estimated at a certain time point. This section proposes an empirical regression-based method that can be used for a continuous mediator. The aim is to determine the mean effect over time. The A_g as A_{gt} are used to estimate the effects at each time t under the assumption of time sequence $A_{gt} \rightarrow M_{gt} \rightarrow Y_{gt}$. For analysis of continuous variables, this study used two-way fixed effects (2FE) regression models, to enable simultaneous adjustment for unobserved unit-specific and time-specific confounders [39]. Here we assume that there is no long-term effects, e.g., insomnia status in a given month was assumed to be affected only by the national COVID-19 outbreak and by mobility restriction in that month; similarly, insomnia status in a given month was assumed to be affected only by the severity of the national COVID-19 outbreak and mobility restriction in that month. Under assumptions of linearity, additivity (no interaction), and absence of long-term effects, the estimation of panel regression is a generalized form of DID estimator with multiple treatment level and time-points. According to the literature, the two methods perform equally when both the exposure and mediator are binary with two time-points [39-41].

Three models of Y_{gt} and M_{gt} were built for estimation. First, the role of exposure A_{gt} on outcome Y_{gt} was modeled. Assuming that the treatment effect is additive and that the error

terms are additively separable, the resulting 2FE regression model (Model 1) can be written as

$$E[Y_{gt}|A_{gt} = a_{gt}] = \gamma_A a_{gt} + \gamma_t + \gamma_g \quad (\text{Model 1})$$

In Model 1, γ_g captures the impact of any unobserved but temporally stable characteristic of group g on outcome Y_{gt} ; γ_t captures the impact of time t , which is stable over groups, on outcome Y_{gt} . Additionally, since time has no effect at baseline, Y_{g0} can be viewed as γ_g . Using Y_{g0} to represent the effect of temporally stable characteristics of group g on outcomes Y_{gt} is intuitive. Coefficient, γ_A , which represents the causal effect of A_{gt} on Y_{gt} (i.e., TE), is assumed to be invariant over time.

A similar approach can be used to build a model of the effect of Y_{gt} on both A_{gt} and M_{gt} and to build a model of the effect of M_{gt} on A_{gt} as follows:

$$E[Y_{gt}|A_{gt} = a_{gt}, M_{gt} = m_{gt}] = \theta_A a_{gt} + \theta_M m_{gt} + \theta_t + \theta_g \quad (\text{Model 2})$$

$$E[M_{gt}|A_{gt} = a_{gt}] = \beta_A a_{gt} + \beta_t + \beta_g \quad (\text{Model 3})$$

Model 1 can be used to quantify TE based on γ_A . According to models 1 and 2, the estimated CDE is θ_A , and the estimated PE is $\gamma_A - \theta_A$, which coincides with the difference method. According to models 2 and 3, the estimated NDE is θ_A , and the estimated NIE is $\theta_M \beta_A$. Moreover, when models 1-3 are all correct, $\gamma_A - \theta_A = \theta_M \beta_A$ (see Appendix E for detailed proof), which are the same results obtained by difference method and product method obtains the same estimation results in linear models [42]. The above estimations are applicable when A_{gt} can be binary or continuous. Notably, the results obtained when M_{gt} is binary are identical to those obtained by the DID estimator version mentioned above.

Illustration

Data Description

The method proposed in this study was used to investigate the causal effect of the national COVID-19 outbreak on mental health status and the mediating role of mobility restriction in this causal relationship. The dataset used in this study was for 40 countries throughout the world for the period from March, 2020, to February, 2021. Exposure A_{gt} was the number of deaths per day (monthly average) due to COVID-19. Exposure A_{gt} was used as an indicator of the severity of the national COVID-19 outbreak within a country during this period. Mediator M_{gt} was cellular phone usage in residential areas, which was obtained from Google. Google obtains aggregated and anonymized cellular phone location data from users who have a Google account on their cellular phone and who opted to provide their location data to Google Location History. Cellular phone usage was recorded for six categories of places: workplace, retailers, transit stations, grocery stores, parks, and residences. In this study, residential cellular

phone usage was used as a surrogate of mobility restriction for each country during the period from March, 2020, to February, 2021.

Outcome variable Y_{gt} was the volume of Google searches for “insomnia” during the period March, 2020, to February, 2021, according to Google Trends data. Google search volume was used as a surrogate for mental health status in 40 countries. Accurate translation of ‘insomnia’ into the local languages of the 40 countries was confirmed by using Google Translate for forward translations from Chinese and English and back translation to Chinese and English. For the 40 countries, average monthly values for each variable were collected over a 12-month period (Appendix F). Table S1 in Appendix G provides the descriptive statistics for these variables. The pretreatment time was the month of January, 2020, in which the impacts of the national COVID-19 outbreak had not yet occurred.

Data analysis was demonstrated in two cases (DID design and empirical 2FE regression-based estimation) as mentioned in the estimation section. In the first case, DID estimators were used to quantify the effects of the June, 2020, national COVID-19 outbreak on mobility restriction and insomnia status in July, 2021. The threshold for dichotomizing variable A_g was the monthly median number of deaths in each country. If $A_g = 1$, monthly deaths have increased and the severity of the national COVID-19 outbreak is classified as high. In contrast, if $A_g = 0$, the severity of the pandemic is relatively low. The M_{gt} is also dichotomized by the median value, where $M_{gt} = 1$ suggests that the time people spend in their homes is longer than usual. Standard errors were derived by bootstrap method. In the second case of the data analysis, we estimate the effects of the COVID-19 outbreak on mobility restriction and insomnia status in the period from March, 2020, to February, 2021, and all variables were considered continuous. Appendix H presents the models of the volumes of ‘insomnia’ searches and the model of the extent of mobility restriction. Since fixed time effects were controlled and eliminated during the estimation process, all 12 months of data were used in the analysis. In this estimation, standard deviation was used to standardize the number of deaths. Delta method was used to estimate standard error. All analyses were performed with R software, version 3.6.3 (R Foundation for Statistical Computing).

DID estimation

Table 1 presents the TE, CDE, PE, NDE and NIE. The TE of the severity of the national COVID-19 outbreak had a significant positive effect on the tendency of insomnia (10.16, 95% CI 2.81 to 17.51). This indicated that the tendency of insomnia increased during the COVID-19 pandemic outbreak, which is consistent with the literature. Additionally, the effect of A on M (DID_{M_{gt},A_g}) was 0.59 (95% CI 0.35 to 0.83) while the effect of M on Y ($DID_{Y_{gt},M_{gt}|A_g=1}$)

was 7.73 (95% CI 2.46 to 13.00), which respectively indicated that (1) mobility restrictions increased as the severity of the national COVID-19 outbreak increased and that (2) the tendency of insomnia increased as mobility restrictions increased. Mediation analysis revealed a significant NIE (4.62, 95% CI 1.27 to 7.97, with Proportion mediated as 0.45), i.e., that mobility restriction mediated the causal effects of COVID-19-induced insomnia and explained approximately 45 % of the causal mechanism. Since 55% of the effect was not explained by the mediation pathway, other important mechanisms may have had roles and need further study.

Empirical Regression-based Estimation

This study used 2FE models to estimate the effect over the year. Table S2 (see Appendix I) shows the maximum likelihood estimates for the coefficients in Models 4-6. Table 2 presents the estimated effects (TE, CDE, PE, NDE and NIE). All effects were positive. However, only NIE was statistically significant (0.41, 95% CI = 0.18 (0.06, 0.76)) with the proportion mediated equal to 0.29, which provided evidence that the mediating effect of mobility restriction on the association between the severity of the national COVID-19 outbreak and the tendency of insomnia continued throughout the entire 1-year period of the analysis.

Discussion

To our best knowledge, this study is the first to extend causal mediation analysis to DID design and panel data settings. This study provided formal counterfactual definitions for direct and indirect effects in panel data settings and DID design, including both the identification and required assumptions. Researchers can use general statistical software to implement the proposed DID estimators and regression-based estimators.

Several limitations of the proposed methods are worth noting. First, the empirical regression-based estimator and model require strong causal assumptions. For example, the model assumes the absence of interactions between variables, including interactions between time and group, between time and variables, and between group and variables. However, interactions may exist in many real-world scenarios. Additionally, non-linear outcome models are widely used in epidemiology research. Additional studies are needed to develop a general methodology that is applicable under widely varying conditions. Moreover, our analysis assumed time sequences between variables, which must be confirmed by substantial background knowledge when conducting data analysis. Sensitivity analysis techniques must be developed to assess bias when assumptions are violated. Finally, all analyses in this study are

based on ecological data and the causal inference is only applicable in ecological level.
Inferring causality to individual level will be subject to ecological fallacy.

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Table 1. Estimations for all effects of the covid-19 severities on search volumes of “insomnia”.

	Estimation	SD	95%CI	P value
Total effect	10.16	3.75	(2.81, 17.51)	0.006
Controlled direct effect	6.67	4.49	(-2.13, 15.47)	0.13
Portion Eliminated	3.49	3.75	(-3.86, 10.84)	0.34
Effect of A on M (DID_{M_{gt},A_g})	0.59	0.12	(0.35, 0.83)	<0.001
Effect of M on Y ($DID_{Y_{gt},M_{gt} A_g=1}$)	7.73	2.69	(2.46, 13.00)	0.004
Natural direct effect	5.54	2.89	(-0.12, 11.20)	0.053
Natural indirect effect	4.62	1.71	(1.27, 7.97)	0.013
Proportion eliminated	0.34			
Proportion mediated	0.45			

Table 2. Estimation for all effects of the covid-19 severities on search volumes of “insomnia”.

	Estimation	SD	95%CI	P value
Total effect	1.41	0.72	(-0.004, 2.82)	0.051
Controlled direct effect =Natural direct effect	1.00	0.74	(-0.46, 2.45)	0.17
Portion Eliminated =Natural indirect effect	0.41	0.18	(0.06, 0.76)	0.021
Proportion Mediated = Proportion Eliminated	0.29			