
Causal Attribution Analysis for Continuous Outcomes

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Abstract

Previous studies have extensively addressed the attribution problem for binary outcome variables. However, in many practical scenarios, the outcome variable is continuous, and simply binarizing it may result in information loss or biased conclusions. To address this issue, we propose a series of posterior causal estimands for retrospectively evaluating multiple correlated causes from a continuous outcome. These estimands include posterior intervention effects, posterior total causal effects, and posterior natural direct effects. Under assumptions of sequential ignorability, monotonicity, and perfect positive rank, we show that the posterior causal estimands of interest are identifiable and present the corresponding identification equations. We also provide a simple but effective estimation procedure and establish asymptotic properties of the proposed estimators. An artificial hypertension example and a real developmental toxicity dataset are employed to illustrate our method.

1. Introduction

In social science (VanderWeele, 2012), health risk assessment (Khoury et al., 2004), legal contexts (Sanders et al., 2021), and explainable artificial intelligence (Galhotra et al., 2021), researchers are interested not only in assessing the effects of causes (Rosenbaum & Rubin, 1983; Robins et al., 1994; Angrist et al., 1996; Bang & Robins, 2005; Ding et al., 2011; Zhao et al., 2012; Jiang et al., 2016; Fan Li & Zaslavsky, 2018; Yang et al., 2020), but also in inferring causes from specific outcomes (Pearl, 1995; Dawid, 2000; Tian & Pearl, 2000; Kuroki & Cai, 2011; Dawid & Musio, 2022; Pearl, 2022; Lu et al., 2023; Li et al., 2023). For instance, for hypertensive patients, researchers may retrospectively eval-

uate whether the development of hypertension was caused by dietary habits, exercise routines, and physical characteristics. Additionally, such questions are also very common in developmental toxicity risk studies, where researchers in clinical trials aim to determine whether abnormal weight loss in pups is caused by potentially toxic agents, organ disease, or other risk factors.

To explain such retrospective studies, researchers need to use counterfactual scenarios that imagine what would have happened if certain conditions experienced previously had been different (Lu et al., 2023). For example, for a patient with high blood pressure who never exercises, it is inferred what his blood pressure would have been like if there had been an intervention to make him exercise regularly. Similarly, for a pup exposed to a toxic reagent and abnormally thin, it is inferred how its body weight would have changed if it had received a placebo instead. Pearl (2000) introduced a three layer causal hierarchy, comprising association, intervention, and counterfactual levels. Retrospective causal analysis, which assesses the causes of observed effects, falls under the third level of this framework. The first two levels primarily involve predicting or evaluating the effects of interventions. In contrast, the third level focuses on determining whether observed outcomes can be attributed to prior interventions or exposures (Pearl, 2015; Dawid et al., 2014; 2015).

While randomized experiments and standard assumptions effectively address the first two levels of causation (Rosenbaum & Rubin, 1983; Pearl, 2014), they are not enough to address the challenges posed by the third level (Dawid & Musio, 2022). This limitation presents a significant challenge for traditional causal inference methods when dealing with such retrospective analysis problems. To formally answer such questions, Pearl (1999) outlined three counterfactual definitions of causal relationships to capture the necessity or sufficiency of a cause for a given binary effect. Additionally, Dawid et al. (2014) introduced the probability of causation to infer the cause for a given binary effect. When there are multiple potentially correlated causes, Lu et al. (2023) and Li et al. (2023) introduced posterior causal effects under observed post-treatment variables to retrospectively deduce causes from a single effect and multiple effects, respectively. In many clinical trials, the outcome variables of interest may be continuous, such as weight,

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blood pressure, and income. However, most existing literature primarily conducts attribution analysis for binary outcomes (Pearl, 1999; Dawid et al., 2014; Lu et al., 2023). Research on continuous outcome variables remains limited, and the formal definitions, identification expressions, and estimation procedures for continuous outcomes require further exploration.

In this paper, we define posterior intervention effects, posterior total causal effects, posterior natural direct effects, and posterior natural indirect effects based on an event related to the continuous outcome variable. We use these definitions to assess multiple correlated causes of a given continuous effect. Interestingly, we establish that under commonly used assumptions of sequential ignorability and perfect positive rank (Heckman et al., 1997), individual treatment effects and posterior intervention causal effects are identifiable. Furthermore, assuming monotonicity among multiple causes (Lu et al., 2023; Li et al., 2023), we demonstrate the identifiability of other posterior causal estimands and provide their identification equations. We also present simplified identification results for the proposed causal estimands under a known directed acyclic graph. To estimate these causal estimands, we propose a two-step approach: first estimating counterfactual mappings and individual treatment effects for each unit, followed by estimating other posterior causal estimands. We also illustrate the application of these posterior causal estimands using an artificial hypertension example. Proofs of all theoretical results are provided in the supplementary materials.

This paper is structured as follows. Section 2 presents the notation and definitions. Section 3 discusses the identifiability of the proposed posterior causal estimands. Section 4 outlines a two-step estimation method for the proposed estimands. In Section 5, we employ an artificial hypertension example to illustrate our proposed method. Finally, Section 6 concludes with a brief summary.

2. Notation and definitions

Assuming that we observe n independent and identically distributed samples from a superpopulation. We first consider the scenario with a single cause and a single outcome. For each unit i , let X_i represent a binary potential cause, where $X_i = 1$ indicates receiving treatment, and $X_i = 0$ indicates receiving control. Let Y_i be the observed continuous outcome. Let $Y_{i,X=0}$ and $Y_{i,X=1}$ denote the potential outcomes corresponding to $X_i = 0$ and $X_i = 1$, respectively. Many common measurements, such as weight, blood pressure, and income, are typically continuous, but Y_i may fall within a specific interval of interest denoted as \mathcal{E}_i . Therefore, given the evidence (X_i, \mathcal{E}_i) , we aim to evaluate the effect of changes in X_i on the outcome Y_i , thereby evaluating the likelihood of X_i being the cause of event \mathcal{E}_i .

Next, we consider the case with multiple causes $X = (X_1, \dots, X_p)$ and a single outcome Y , where X is a binary vector of causes, and the causes may affect each other. Without loss of generality, we assume that the causes are arranged in a topological order such that X_l is not a cause of X_k for $k < l$. For example, X is a sequence of observations ordered in time, or X consists of variables in a directed acyclic graph (DAG) where X_k is not a descendant of X_l for $k < l$. For generic sets of variables W and U , we use W_u to denote the potential outcome of W that would have resulted if U were intervened to level u . In particular, if $W = (W_1, \dots, W_s)$, then $W_u = \{(W_1)_u, \dots, (W_s)_u\}$. We make the consistency assumption that connects observed variables to potential outcomes, i.e., $W_u = W$ if $U = u$ (Pearl, 2015). We further suppose the composition assumption holds in the sense that for any variable sets W, V and $U, W_{vu} = W_v$ if $U_v = u$ (Pearl, 2015). The consistency assumption can be viewed as a special case of the composition assumption if V is empty.

To measure how likely X_k is a cause of the continuous effect given observed evidence (x, \mathcal{E}) , we extend the concept of *posterior total causal effect* (postTCE) defined by Lu et al. (2023) as follows:

$$\begin{aligned} \text{PostTCE}(X_k \Rightarrow Y \mid x, \mathcal{E}) \\ = E(Y_{X_k=1} - Y_{X_k=0} \mid x, \mathcal{E}), \end{aligned}$$

where we use “ x ” to represent “ $X = x$ ” for notational simplicity. It is important to note that this definition includes the event \mathcal{E} defined by the observed outcome, and cannot simply be considered as a conditional average causal effect (CATE). As advocated by Lu et al. (2023), a larger value of the posterior total causal effect indicates that the effect or outcome is more attributable to the cause X_k . The cause that produces the largest posterior total causal effect is usually considered the highest risk factor.

Similar to the direct causal effect considered by Pearl (2000), we define the posterior natural direct effect of X_k on Y given the observed evidence (x, \mathcal{E}) , which quantifies the effect of X_k on Y not mediated through intermediate variables. Let $A_k = (X_1, \dots, X_{k-1})$ and $D_k = (X_{k+1}, \dots, X_p)$. Then $X = (A_k, X_k, D_k)$, and $x = (a_k, x_k, d_k)$ denotes a value of X . Given the evidence (x, \mathcal{E}) , the *posterior natural direct effect* (postNDE) of X_k on Y is:

$$\begin{aligned} \text{PostNDE}(X_k \Rightarrow Y \mid x, \mathcal{E}) \\ = E\{Y_{X_k=1, D_k(a_k, 0)} - Y_{X_k=0} \mid x, \mathcal{E}\}, \end{aligned}$$

where $D_k(a_k, 0)$ is the potential outcome under $(A_k, X_k) = (a_k, 0)$. Throughout this paper, we use $D_k(a_k, x_k)$ and $(D_k)_{a_k, x_k}$ interchangeably in the nested potential outcomes. The postNDE describes the effect observed when each individual in the subpopulation (x, \mathcal{E}) switches from $X_k = 0$ to $X_k = 1$, while keeping D_k at its value when $X_k = 0$.

Parallel to the natural indirect effect considered by Pearl (2000), we also define the *posterior natural indirect effect* (postNIE) of X_k on Y given the evidence (x, \mathcal{E}) as follows:

$$\begin{aligned} \text{PostNIE}(X_k \Rightarrow Y \mid x, \mathcal{E}) \\ = E\{Y_{X_k=1} - Y_{X_k=1, D_k(a_k, 0)} \mid x, \mathcal{E}\}. \end{aligned}$$

The postNIE quantifies, for each individual in the subpopulation (x, \mathcal{E}) , the effect observed when X_k is set to $X_k = 1$, while all intermediate variables along the pathway from X_k to Y change from state $D_k(a_k, 1)$ to state $D_k(a_k, 0)$. Through the definitions, we have that:

$$\begin{aligned} \text{PostTCE}(X_k \Rightarrow Y \mid x, \mathcal{E}) \\ = \text{PostNDE}(X_k \Rightarrow Y \mid x, \mathcal{E}) + \text{PostNIE}(X_k \Rightarrow Y \mid x, \mathcal{E}). \end{aligned}$$

Given the observed evidence (x, \mathcal{E}) , in addition to assessing the a posteriori total, direct and indirect effects of a particular cause X_k , we need to consider assessing the synergistic effects of a joint intervention with all possible causes in an alternative state $X = x'$. Indeed, synergistic effects are important in many applications. For example, having heart disease alone may have a limited effect on blood pressure, whereas the combination of an unhealthy diet and heart disease may jointly contribute to elevated blood pressure. Therefore, the *posterior intervention causal effect* (postICE) for another state $X = x'$ is defined as follows:

$$\text{PostICE}(Y_{x'} \mid x, \mathcal{E}) = E(Y_{x'} - Y \mid x, \mathcal{E}). \quad (1)$$

The *individual treatment effect* (ITE) for any pair (x', x^*) can be defined as $\text{ITE}(x', x^*) = Y_{x'} - Y_{x^*}$, representing the difference in potential outcomes for each individual under two different treatment conditions. Inferring ITEs presents a fundamental challenge because we can only observe one potential outcome for each unit (Rosenbaum & Rubin, 1983).

3. Identifiability of posterior causal estimands and required assumptions

3.1. Assumptions required for identifiability

Define $W = (X, Y)$ and let $W_{r:s}$ denote a subvector $(W_r, W_{r+1}, \dots, W_s)$ of W for $r \leq s$. Let $w_{r:s}^* = (w_r^*, \dots, w_s^*) \preceq w_{r:s} = (w_r, \dots, w_s)$ denote that $w_i^* \leq w_i$ for all $r \leq i \leq s$. To identify the proposed estimands, we make the following commonly used assumptions in previous studies (Heckman et al., 1997; Pearl, 2000; 2014; Lu et al., 2023; Li et al., 2023).

Assumption 3.1 (Sequential ignorability). We consider the following assumptions:

- (i) there is no confounding between W_s and $W_{1:s-1}$, i.e., $(W_s)_{w_{1:s-1}} \perp\!\!\!\perp W_{1:s-1}$ for all $w_{1:s-1}$ and $s = 2, \dots, p+1$;

- (ii) the elements in $\{(W_s)_{w_{1:s-1}}\}_{s=1}^{p+1}$ are mutually independent for any given $w_{1:p}$.

The independence condition in Assumption 3.1 can be relaxed by introducing the baseline covariates, and we omit it for simplicity. The Assumption 3.1(i) implies that the potential outcome of each variable is independent of the prior variables in causal order. Under the Assumption 3.1(i), if W_s has a nonparametric causal structural model $W_s = m_s(W_{1:s-1}, \epsilon_s)$ with an unknown function $m_s(\cdot, \epsilon_s)$ and a error variable $\epsilon_s \perp\!\!\!\perp W_{1:s-1}$, then Assumption 3.1(ii) holds naturally because Assumption 3.1(i) implies that $\epsilon_s \perp\!\!\!\perp \epsilon_{1:s-1}$ for $s = 2, \dots, p+q$, which further implies Assumption 3.1(ii). Assumption 3.1 rules out unobserved confounders between any two variables in W . However, each variable X_k may still confound the relationship between Y and X_l , or between X_l and X_s , provided $k < l, s$. Assumption 3.1 is frequently employed in causal inference with complex systems, including mediation analysis (Imai et al., 2010) and longitudinal data involving time-dependent confounders (Robins, 2000).

Assumption 3.2 (Monotonicity). For $s = 2, \dots, p$, we assume that $(W_s)_{w_{1:s-1}^*} \leq (W_s)_{w_{1:s-1}}$ whenever $w_{1:s-1}^* \preceq w_{1:s-1}$ holds.

Assumption 3.2 implies that each cause has a non-negative effect on subsequent causes. This assumption is commonly expressed in epidemiology as “no prevention”, meaning that no individual is helped by exposure to a risk factor. To identify the posterior causal estimands, Lu et al. (2023) and Li et al. (2023) also introduce the same monotonicity assumption across multiple potentially correlated causes. The validity of monotonicity cannot be tested directly, but under Assumption 3.1, the monotonicity can be falsified by imposing testable restrictions on the observed data distribution. For example, for any $w_{1:s-1}^* \preceq w_{1:s-1}$, the following equality can be used to falsify the monotonicity assumption:

$$\begin{aligned} \text{pr}(W_s = 1 \mid W_{1:s-1} = w_{1:s-1}^*) \\ \leq \text{pr}(W_s = 1 \mid W_{1:s-1} = w_{1:s-1}). \end{aligned}$$

Assumption 3.3 (Perfect positive rank). We assume that $W_{p+1} = m_{p+1}(W_{1:p}, \epsilon_{p+1})$, or equivalently $Y = m_{p+1}(X, \epsilon_{p+1})$, where ϵ_{p+1} represents a scalar-valued error variable. The unknown link function $m_{p+1}(X, \cdot)$ is continuous and strictly increasing in ϵ_{p+1} .

For a given binary outcome, Pearl (2000) introduced the monotonicity assumption of the outcome variable with respect to a single cause X , denoted as $Y_{X=0} \leq Y_{X=1}$, which is crucial for identifying the probability of necessity or sufficiency under a single cause X . Lu et al. (2023) also considered the monotonicity assumption of the binary outcome with respect to multiple correlated causes X , denoted as

$Y_{X=x^*} \leq Y_{X=x}$ for any $x^* \preceq x$. However, this monotonic relationship may not be applicable when the outcome variable is continuous. Therefore, we introduce Assumption 3.3, which is commonly used to identify individual treatment effects or quantile treatment effects in counterfactual causal inference literature (Heckman et al., 1997; Chernozhukov & Hansen, 2005), but it is a relatively novel assumption in retrospective attribution analysis.

The basic restriction in Assumption 3.3 is also referred to as the rank preservation or rank invariance (Heckman et al., 1997; Chernozhukov & Hansen, 2005; Vuong & Xu, 2017; Feng et al., 2020). If an individual with regular exercise and without heart disease (i.e., $x = (1, 0)$) has the lowest blood pressure in the subpopulation $\{(X_i, Y_i) : X_i = x\}$, then according to Assumption 3.3, an individual with no exercise and heart disease (i.e., $x' = (0, 1)$) should also have the lowest blood pressure in the corresponding subpopulation $\{(X_i, Y_i) : X_i = x'\}$; and vice versa. The strict monotonic increase of ϵ can be changed to the strict monotonic decrease without affecting the subsequent discussion. For simplicity, we assume the strict monotonic increase. For any given error ϵ_{p+1}^* , Assumption 3.3 requires that the relative rank or quantile of $Y_x \equiv m_{p+1}(x, \epsilon_{p+1}^*)$ be the same as that of $Y_{x'} \equiv m_{p+1}(x', \epsilon_{p+1}^*)$ for any $x \neq x'$. A stronger version of Assumption 3.3 assumes that the error term ϵ_{p+1} is additive, that is, $Y = m_{p+1}^*(X) + \epsilon_{p+1}$ for some real-valued function $m_{p+1}^*(\cdot)$. Moreover, the heteroscedasticity model also satisfies Assumption 3.3: $Y = m_{p+1}^*(X) + \sigma(X)\epsilon_{p+1}$, for some real-valued function $m_{p+1}^*(X)$ and positive function $\sigma(X)$.

3.2. Identification equations of posterior causal estimands

Under Assumptions 3.1 and 3.3, we first consider the identifiability of ITEs and postICEs. Let S_{Y_x} denote the support of the potential outcome Y_x , which can be identified by $S_{Y|X=x}$ under Assumption 3.1. The key to our identification strategy is to match the potential outcome $Y_x \equiv m_{p+1}(x, \epsilon_{p+1})$ with another potential outcome $Y_{x'} \equiv m_{p+1}(x', \epsilon_{p+1})$ through a mapping $\phi_{x \rightarrow x'}(\cdot)$, such that $Y_{x'} = \phi_{x \rightarrow x'}(Y_x)$. This mapping $\phi_{x \rightarrow x'}(\cdot)$ is termed a counterfactual mapping (Vuong & Xu, 2017; Feng et al., 2020), because it allows us to find the counterfactual outcome $Y_{x'}$ from Y_x using the function $\phi_{x \rightarrow x'}(\cdot)$, and vice versa.

Let $m_{p+1}^{-1}(x, \cdot)$ be the inverse function of $m_{p+1}(x, \cdot)$. According to Assumption 3.3, for any pair $(X, Y) = (x, y)$, we can uniquely represent the error term as $\epsilon_{p+1} = m_{p+1}^{-1}(x, y)$, although the specific form of $m_{p+1}^{-1}(x, y)$ is unknown. Therefore, $Y_{x'}$ is uniquely defined by $\phi_{x \rightarrow x'}(y) \equiv m_{p+1}\{x', m_{p+1}^{-1}(x, y)\}$ for each $y \in S_{Y_x}$. Moreover, the counterfactual mapping $\phi_{x \rightarrow x'}(\cdot)$ is a continuous and

strictly increasing function from S_{Y_x} onto $S_{Y_{x'}}$. Thus, if we can identify $\phi_{x \rightarrow x'}(y)$ for all $y \in S_{Y_x}$ and $x \neq x'$, we can recover ITEs for each individual.

Lemma 3.4. *Under Assumptions 3.1 and 3.3, for any $y \in S_{Y_x}$, the counterfactual mapping $\phi_{x \rightarrow x'}(\cdot)$ is identified by the continuous extension of*

$$\phi_{x \rightarrow x'}(y) = F_{x'}^{-1}\{F_x(y)\}, \quad \forall y \in S_{Y_x}^\circ, \quad (2)$$

where $F_x(y) = \text{pr}(Y \leq y | X = x)$ and $S_{Y_x}^\circ$ is the interior of S_{Y_x} . Moreover, the ITEs of every individual in the population can be identified.

Lemma 3.4 establishes the identifiability of the counterfactual mapping $\phi_{x \rightarrow x'}(\cdot)$ on S_{Y_x} constructively by matching the quantiles of Y_x and $Y_{x'}$. We provide further intuition for Lemma 3.4. When we observe an individual in the subpopulation $\{(X_i = x, Y_i)\}$ with the highest blood pressure, we can recover the joint distribution of individual i by identifying the individual with the highest blood pressure in each observed subgroup $\{(X_j = x', Y_j)\}$ for any $x' \neq x$. Given the identifiability of the counterfactual mapping and ITEs, the postICEs can also be identified using an inverse probability weighting expression (Horvitz & Thompson, 1952),

$$\text{PostICE}(Y_{x'} | x, \mathcal{E}) = E \left[\frac{\mathbb{I}(X = x, \mathcal{E})}{\text{pr}(X = x, \mathcal{E})} \{ \phi_{x \rightarrow x'}(Y) - Y \} \right], \quad (3)$$

where $\mathbb{I}(\cdot)$ denotes the indicator function.

Assumptions 3.1 and 3.3 establish the identifiability of ITEs and postICEs. However, they are not sufficient to ensure the identifiability of other posterior causal estimands (e.g., postNDEs and PostNIEs) when considering a specific cause X_k . This is due to the challenge posed by identifying expectations of the nested potential outcomes given the observed evidence. For example, the postNDE involves the conditional expectation of the cross-world intervention potential outcome $Y_{X_k=1, D_k(a_k, 0)}$, which cannot be identified using Lemma 3.4 alone. Before formally identifying these expectations, we provide another lemma for identifying the conditional probability of the counterfactual outcomes of the causes.

Lemma 3.5. *Under Assumptions 3.1 and 3.2, given the observed evidence $(a_k, x_k, d_k, \mathcal{E})$, let $d_k^* = (x_{k+1}^*, \dots, x_p^*)$ and $d_k = (x_{k+1}, \dots, x_p)$.*

(i) *For $d_k^* \preceq d_k$, we have,*

$$\begin{aligned} \text{pr}\{D_k(a_k, 0) = d_k^* | a_k, 1, d_k\} \\ = \prod_{s=k+1}^p \{(1 - x_s^*) + (2x_s^* - 1)x_s R_{0s}\}, \end{aligned}$$

$$\text{where } R_{0s} = \frac{\text{pr}(X_s = 1 | a_k, 0, x_{k+1}^*, \dots, x_{s-1}^*)}{\text{pr}(X_s = 1 | a_k, 1, x_{k+1}, \dots, x_{s-1})}.$$

(ii) For $d_k \preceq d_k^*$, we have,

$$\begin{aligned} \text{pr}\{D_k(a_k, 1) = d_k^* \mid a_k, 0, d_k\} \\ = \prod_{s=k+1}^p \{x_s^* + (1 - 2x_s^*)(1 - x_s)R_{1s}\}, \\ \text{where } R_{1s} = \frac{\text{pr}(X_s = 0 \mid a_k, 1, x_{k+1}^*, \dots, x_{s-1}^*)}{\text{pr}(X_s = 0 \mid a_k, 0, x_{k+1}, \dots, x_{s-1})}. \end{aligned}$$

Theorem 3.6. Under Assumptions 3.1 and 3.2, given the evidence $(a_k, x_k, d_k, \mathcal{E})$, the postNDE, postNIE, and postTCE of X_k on Y can be identified as follows:

(i) given $x_k = 1$, for any $x_k^* \in \{0, 1\}$, we have,

$$\begin{aligned} E\{Y_{x_k^*, D_k(a_k, 1)} \mid x, \mathcal{E}\} &= E(Y_{x_k^*, d_k} \mid x, \mathcal{E}), \\ E\{Y_{x_k^*, D_k(a_k, 0)} \mid x, \mathcal{E}\} \\ &= \sum_{d_k^* \preceq d_k} E(Y_{x_k^*, d_k^*} \mid x, \mathcal{E}) \text{pr}\{D_k(a_k, 0) = d_k^* \mid x\}, \end{aligned}$$

(ii) given $x_k = 0$, for any $x_k^* \in \{0, 1\}$, we have,

$$\begin{aligned} E\{Y_{x_k^*, D_k(a_k, 0)} \mid x, \mathcal{E}\} &= E(Y_{x_k^*, d_k} \mid x, \mathcal{E}), \\ E\{Y_{x_k^*, D_k(a_k, 1)} \mid x, \mathcal{E}\} \\ &= \sum_{d_k \preceq d_k^*} E(Y_{x_k^*, d_k^*} \mid x, \mathcal{E}) \text{pr}\{D_k(a_k, 1) = d_k^* \mid x\}, \end{aligned}$$

where $E(Y_{x_k^*, d_k^*} \mid x, \mathcal{E})$ can be identified by Lemma 3.4, and $\text{pr}\{D_k(a_k, x_k') = d_k^* \mid x\}$ for $x_k' \in \{0, 1\}$ can be identified by Lemma 3.5.

Theorem 3.6 illustrates that with the additional monotonicity assumption 3.2, we can also identify postNDE, postNIE, and postTCE. In some cases, we may only want to conduct attribution analysis for continuous effects based on evidence from a subset of (X, \mathcal{E}) , while Lemma 3.4 and Theorem 3.6 are based on fully observed evidence (x, \mathcal{E}) . For some $s < p$, let $(X', \mathcal{E}) = (X_{i_1}, \dots, X_{i_s}, \mathcal{E})$ denote a subset of (X, \mathcal{E}) . We can summarize the remaining set $X \setminus X'$ to obtain the expected results for the subset (X', \mathcal{E}) based on Lemma 3.4 and Theorem 3.6. To simplify the exposition, we omit this part. We also refer to Corollary 1 in Lu et al. (2023) and Theorem 3 in Li et al. (2023) for parallel results.

3.3. Identification equations of posterior causal effects under causal networks

In this section, we consider the causal structure of observed variables (X, Y) represented by a directed acyclic graph (DAG) or network. We aim to present the simplified identification expressions for the posterior causal estimands given a known DAG. For $k = 1, \dots, p$, let Pa_k and Pa_Y denote the sets of parents of X_k and Y in the graph, respectively. The joint probability distribution $\text{pr}(X, Y)$ can be factorized as $\text{pr}(X, Y) = \prod_{k=1}^p \text{pr}(X_k \mid \text{Pa}_k) \text{pr}(Y \mid \text{Pa}_Y)$. We assume $\text{pr}(x, y) > 0$ for each (x, y) . For a given graph,

the sequential ignorability assumption (Assumption 3.1) posits that there is no unobserved variable intervening between any two or more nodes in the DAG. The monotonicity assumption 3.2 implies that each node X_k has a positive individual monotonic effect on its child nodes. Furthermore, Assumption 3.3 can be simplified such that the unknown link function $m_{p+1}(\text{Pa}_Y, \epsilon_{p+1})$ only needs to be continuous with respect to Pa_Y and strictly increasing in ϵ_{p+1} . Specifically, the continuous outcome variable is only required to perfectly match the quantiles of potential outcomes under different realizations of its parent nodes Pa_Y . We define a simpler counterfactual mapping $\phi_{\text{Pa}_Y \rightarrow \text{Pa}_Y'}(\cdot)$ to characterize the mapping relationship of the parent nodes of Y under different realizations Pa_Y and Pa_Y' . The theoretic results presented in Lemma 3.4 and (3) can be simplified for a given graph.

Corollary 3.7. Suppose that the causal network of (X_1, \dots, X_p, Y) is a DAG. Then, under Assumptions 3.1 and 3.3, for any $x \neq x'$ and $y \in \mathcal{S}_{Y_x}$, let $\text{Pa}_Y \subset x$ and $\text{Pa}_Y' \subset x'$, the counterfactual mapping $\phi_{x \rightarrow x'}(\cdot)$ can be reduced as follows:

$$\phi_{x \rightarrow x'}(y) = \phi_{\text{Pa}_Y \rightarrow \text{Pa}_Y'}(y) = F_{\text{Pa}_Y}^{-1}\{F_{\text{Pa}_Y'}(y)\}, \quad \forall y \in \mathcal{S}_{Y_x}^\circ,$$

where $m_{\text{Pa}_Y}(y) = \text{pr}(Y \leq y \mid \text{Pa}_Y = \text{Pa}_Y)$. Moreover, the ITEs of every individual in the population can be identified. The following equation also holds:

$$\begin{aligned} E(Y_{x'} \mid x, \mathcal{E}) &= E(Y_{\text{Pa}_Y'} \mid \text{Pa}_Y, \mathcal{E}) \\ &= E\left\{\frac{\mathbb{I}(\text{Pa}_Y = \text{Pa}_Y', \mathcal{E})}{\text{pr}(\text{Pa}_Y = \text{Pa}_Y', \mathcal{E})} \phi_{\text{Pa}_Y \rightarrow \text{Pa}_Y'}(Y)\right\}, \end{aligned}$$

and the postICE is identifiable.

Corollary 3.7 indicates that the counterfactual mapping depends only on Pa_Y , the postICEs and ITEs can be calculated using the low-dimensional probabilities $\text{pr}(Y \leq y \mid \text{Pa}_Y)$. Similarly, the identification equations of the conditional probability $\text{pr}\{D_k(a_k, 0) = d_k^* \mid x\}$ can also be simplified by replacing R_{0s} and R_{1s} in Lemma 3.5 with R_{0s}^* and R_{1s}^* :

$$\begin{aligned} R_{0s}^* &= \text{pr}(X_s = 0 \mid \text{Pa}_s^*) / \text{pr}(X_s = 0 \mid \text{Pa}_s), \\ R_{1s}^* &= \text{pr}(X_s = 1 \mid \text{Pa}_s^*) / \text{pr}(X_s = 1 \mid \text{Pa}_s), \end{aligned} \quad (4)$$

where $\text{Pa}_s^* \subset (a_k, 0, x_{k+1}^*, \dots, x_{s-1}^*)$ and $\text{Pa}_s \subset (a_k, 1, x_{k+1}, \dots, x_{s-1})$. The following Corollary 3.8 provides a simplified identification equation of other posterior causal estimands for a given graph.

Corollary 3.8. Suppose that the causal network of (X_1, \dots, X_p, Y) is a DAG. Under Assumptions 3.1 and 3.2, given the observed evidence $(a_k, x_k, d_k, \mathcal{E})$, the postNDE, postNIE, and postTCE of X_k on Y can be identified using the following equations:

(i) when $x_k = 1$, for any $x_k^* \in \{0, 1\}$, we have,

$$\begin{aligned} E\{Y_{x_k^*, D_k(a_k, 1)} \mid x, \mathcal{E}\} &= E(Y_{\text{pa}^*} \mid \text{pa}_y, \mathcal{E}), \\ E\{Y_{x_k^*, D_k(a_k, 0)} \mid x, \mathcal{E}\} &= \\ \sum_{d_k^* \leq d_k} E(Y_{\text{pa}_y^*} \mid \text{pa}_y, \mathcal{E}) \text{pr}\{D_k(a_k, 0) = d_k^* \mid x\}, \end{aligned}$$

(ii) when $x_k = 0$, for any $x_k^* \in \{0, 1\}$, we have,

$$\begin{aligned} E\{Y_{x_k^*, D_k(a_k, 0)} \mid x, \mathcal{E}\} &= E(Y_{\text{pa}^*} \mid \text{pa}_y, \mathcal{E}), \\ E\{Y_{x_k^*, D_k(a_k, 1)} \mid x, \mathcal{E}\} &= \\ \sum_{d_k \leq d_k^*} E(Y_{\text{pa}_y^*} \mid \text{pa}_y, \mathcal{E}) \text{pr}\{D_k(a_k, 1) = d_k^* \mid x\}, \end{aligned}$$

where $E(Y_{\text{pa}_y^*} \mid \text{pa}_y, \mathcal{E})$ and $E(Y_{\text{pa}^*} \mid \text{pa}_y, \mathcal{E})$ can be identified by Corollary 3.7 for $\text{pa}_y^* \subset (a_k, x_k^*, d_k)$ and $\text{pa}_y \subset (a_k, x_k^*, d_k^*)$, and $\text{pr}\{D_k(a_k, x_k') = d_k^* \mid x\}$ can be identified by Lemma 3.5 and (4) for any $x_k' \in \{0, 1\}$.

The theoretic results in this section are very useful in practice and can greatly simplify the computation, especially when the dimensionality of the causes is large. We suggest that practitioners first obtain a simple DAG, or a larger one but with the correct order of nodes, from expert knowledge or conditional independence tests.

4. Estimation

In this section, we propose a simple but effective method for estimating the counterfactual outcome mapping $\phi_{x \rightarrow x'}(\cdot)$ as well as the posterior causal estimands. Let $\{(X_i, Y_i) : i = 1, \dots, n\}$ be the independent and identically distributed samples generated according to Assumptions 3.1-3.3. Our estimation procedure consists of two steps: first, for each observation $(X_i, Y_i) = (x, y)$, we estimate the counterfactual mapping $\phi_{x \rightarrow x'}(y)$ for $x' \neq x$ using a simple estimator that minimizes a convex population objective function and constructs pseudo samples of the counterfactual outcomes for all individuals. In the second step, we nonparametrically estimate the posterior causal estimands based on the counterfactual mapping $\phi_{x \rightarrow x'}(\cdot)$. For simplicity, let \mathcal{S}_{Y_x} denote a compact interval $[y_x^l, y_x^u]$ for any $X = x$, where $-\infty < y_x^l < y_x^u < +\infty$. We also assume the compact support $[y_x^l, y_x^u]$ is known. Otherwise, it can be estimated using the methods proposed in Korostelev & Tsybakov (2012). To establish the asymptotic properties of the estimators to be proposed in this section, we make the following assumptions.

Assumption 4.1. (i) The function $m_{p+1}(x, \epsilon_{p+1})$ is continuously differentiable in error term ϵ_{p+1} for any $X = x$; (ii) The probability density function of the error term ϵ_{p+1} is continuous; (iii) $\inf_{y \in [y_x^l, y_x^u]} g_x(y) > 0$ for any $X = x$, where $g_x(y) = \partial F_x(y) / \partial y$.

Assumptions 4.1(i) and 4.1(ii) are regularity conditions and, together with Assumptions 3.1 and 3.3, ensure that

the marginal distribution $F_x(y)$ is absolutely continuous with respect to the Lebesgue measure, and its probability density function $g_x(y)$ is also continuous for any $X = x$. Assumption 4.1(iii) is introduced for the sake of simplicity in explanation. Trimming techniques can be employed to relax this assumption, but they may introduce technical complexities.

We now aim to derive a counterfactual outcome mapping $\phi_{x \rightarrow x'}(\cdot)$ from two marginal distributions $F_x(\cdot)$ and $F_{x'}(\cdot)$. For a given $y \in \mathbb{R}$, we define the objective function as follows: $\rho_{x \rightarrow x'}(t; y) = E\{\text{sign}(Y - y) \mid X = x\} \times t - E(|Y - t| \mid X = x')$, where $\text{sign}(u) \equiv 2 \times \mathbb{I}(u > 0) - 1$. The above objective function is motivated by the quantile regression method in Koenker & Bassett (1978) and Feng et al. (2020). Simple calculations reveal that the first-order and second-order conditions of this objective function are given by:

$$\begin{aligned} \frac{\partial \rho_{x \rightarrow x'}(t; y)}{\partial t} &= 2\{F_{x'}(t) - F_x(y)\} = 0, \\ \frac{\partial^2 \rho_{x \rightarrow x'}(t; y)}{\partial t^2} &= 2g_{x'}(t) \geq 0. \end{aligned}$$

The following lemma demonstrates that for any $y \in \mathcal{S}_{Y_x}^\circ$, the objective function $\rho_{x \rightarrow x'}(\cdot; y)$ is uniquely minimized at the counterfactual outcome $\phi_{x \rightarrow x'}(y)$.

Lemma 4.2. Under Assumptions 3.1 and 3.2, $\rho_{x \rightarrow x'}(\cdot; y)$ is continuously differentiable and weakly convex on \mathbb{R} . Additionally, $\rho_{x \rightarrow x'}(\cdot; y)$ is strictly convex on $\mathcal{S}_{Y_x}^\circ$ and uniquely minimized on \mathbb{R} at $\phi_{x \rightarrow x'}(y)$ when $y \in \mathcal{S}_{Y_x}^\circ$.

Lemma 4.2 provides the foundation for our nonparametric estimation of the counterfactual mappings and posterior causal estimands. For the i th observational unit $(X_i = x, Y_i)$, let

$$\begin{aligned} \hat{\rho}_{x \rightarrow x'}(t; Y_i) &= \frac{\sum_{j=1}^n \text{sign}(Y_j - Y_i) \times \mathbb{I}(X_j = x)}{\sum_{j=1}^n \mathbb{I}(X_j = x)} \times t \\ &\quad - \frac{\sum_{j=1}^n |Y_j - t| \times \mathbb{I}(X_j = x')}{\sum_{j=1}^n \mathbb{I}(X_j = x')}. \end{aligned}$$

Hence, we can estimate the counterfactual outcome of the i th unit by,

$$\hat{\phi}_{x \rightarrow x'}(Y_i) = \arg \min_{t \in [y_{x'}^l, y_{x'}^u]} \hat{\rho}_{x \rightarrow x'}(t; Y_i), \text{ if } X_i = x. \quad (5)$$

By separately minimizing $\hat{\rho}_{x \rightarrow x'}(\cdot; Y_i)$ for each unit i with $(X_i = x, Y_i)$, we can estimate the counterfactual outcome for an individual i under state $X = x'$ by the counterfactual mapping $\hat{\phi}_{x \rightarrow x'}(Y_i)$ in (5), and the individual treatment effect $\text{ITE}(x', x^*)$ can be estimated as follows: $\hat{\Delta}_{\text{ITE}, i}(x', x^*) = \hat{\phi}_{x \rightarrow x'}(Y_i) - \hat{\phi}_{x \rightarrow x^*}(Y_i)$. Moreover, other posterior causal estimands can also be constructed using similar moment estimators.

Table 1: Results of posterior intervention causal effects based on different evidence.

PostICE($Y_{x'} \mid x, Y > 140$)	$(x_1, x_4) = (0, 0)$	$(x_1, x_4) = (0, 1)$	$(x_1, x_4) = (1, 0)$	$(x_1, x_4) = (1, 1)$
$(x'_1, x'_4) = (0, 0)$	0.00	-12.43	-1.76	-18.19
$(x'_1, x'_4) = (0, 1)$	3.34	0.00	2.34	-5.43
$(x'_1, x'_4) = (1, 0)$	1.75	-10.01	0.00	-15.75
$(x'_1, x'_4) = (1, 1)$	12.05	5.57	10.86	0.00

5. Example: risk factors for hypertension

In this section, we will use the example provided in Lu et al. (2023) to explain our proposed causal attribution estimands, and assess the posterior causal effects of risk factors on continuous hypertension. Figure 1 presents the causal network and the corresponding conditional probabilities, where Exercise (E), Diet (D), Heart Disease (HD), Heartburn (Hb), and Chest Pain (CP) are potential causes of hypertension. Let $E = 1$ denote no daily exercise, $D = 1$ denote unhealthy diet, $HD = 1$ denote heart disease, $Hb = 1$ denote heartburn, $CP = 1$ denote chest pain, and BP denote continuous blood pressure Y . We generate the data without unobserved confounders to satisfy Assumption 3.1. We also ensure that the relationships among the causes satisfy the monotonicity assumption (Assumption 3.2). For instance, lack of daily exercise ($E = 1$) and poor diet ($D = 1$) are not preventive for heart disease HD . The various potential outcomes also satisfy the perfect rank assumption (Assumption 3.3). Specifically, we modeled blood pressure Y as a continuous outcome variable and ensured that, after binarization, its distribution matches that observed in Lu et al. (2023). The topological order in Figure 1 is $X = (X_1, \dots, X_5) = (E, D, Hb, HD, CP)$. The joint probability of X and Y is calculated by substituting the conditional probabilities from Figure 1 into the following equation: $\text{pr}(E, D, Hb, HD, CP, BP) = \text{pr}(E) \text{pr}(D) \text{pr}(Hb \mid D) \text{pr}(HD \mid E, D) \text{pr}(CP \mid Hb, HD) \text{pr}(BP \mid HD)$. Without loss of generality, we consider the event $\mathcal{E} = \mathbb{I}(Y > 140)$, which indicates whether hypertension is present. The posterior causal estimands can be used to analyze the causes of hypertension for a patient based on the evidence $(x, Y > 140)$.

We first present the postICEs for hypertension, based on different observed evidence, as shown in Table 1. According to Corollary 3.7, we know that the postICEs are only related to the parent nodes of BP , namely X_1 (exercise) and X_4 (heart disease). We find that the largest change in postICEs occurs with the evidence $(x_1, x_4, \mathcal{E}) = (1, 1, Y > 140)$ and the intervened treatment $(x'_1, x'_4) = (0, 0)$. Specifically, for individuals who do not exercise, have heart disease, and have high blood pressure, if they had exercised previously and did not have heart disease, i.e., receiving treatment $(x'_1, x'_4) = (0, 0)$, their blood pressure would significantly decrease by 18.19, that is, $\text{PostICE}(Y_{x'} \mid$

Table 2: Results of marginal probabilities of necessity and posterior causal estimands based on the evidence $\{X = (1, 1, 1, 1, 1), Y > 140\}$.

	X_1	X_2	X_3	X_4	X_5
$\text{PN}^*(X_k \Rightarrow \mathcal{E})$	0.347	0.230	0.133	0.760	0.563
$\text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140)$	0.344	0.207	0	0.722	0
$\text{postNDE}(X_k \Rightarrow Y \mid x, Y > 140)$	3.823	0	0	17.023	0
$\text{postNIE}(X_k \Rightarrow Y \mid x, Y > 140)$	6.805	4.561	0	0	0
$\text{postTCE}(X_k \Rightarrow Y \mid x, Y > 140)$	10.628	4.561	0	17.023	0

$X_1 = 1, X_4 = 1, Y > 140) \approx -18.19$. Conversely, in the evidence $(x_1, x_4, \mathcal{E}) = (0, 0, Y > 140)$, i.e., individuals who exercise, do not have heart disease, and have high blood pressure, if they had not exercised previously and did not have heart disease, i.e., receiving $(x'_1, x'_4) = (1, 0)$, their blood pressure would slightly increase by 1.75, that is, $\text{PostICE}(Y_{x'} \mid X_1 = 0, X_4 = 0, Y > 140) \approx 1.75$.

Since our data generation mechanism ensures the exact same observed data distribution as described in Lu et al. (2023) after binarizing outcome Y , we directly use their posterior total causal effects in their Table 1 for comparative analysis. We use the symbol asterisks (*) to differentiate from the definitions provided in this paper. Specifically, let $Y_{X_k=x_k}^*$ be the binary event $\mathbb{I}(Y_{X_k=x_k} > 140)$, we adopt the definitions of the binary outcomes from Lu et al. (2023), denoted as $\text{PN}^*(X_k \Rightarrow \mathcal{E})$ and $\text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140)$, as follows: $\text{PN}^*(X_k \Rightarrow \mathcal{E}) = \text{pr}(Y_{X_k=0}^* = 0 \mid a_k, X_k = 1, d_k, Y > 140)$,

$$\begin{aligned} \text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140) \\ = E(Y_{X_k=1}^* - Y_{X_k=0}^* \mid x, Y > 140). \end{aligned}$$

The above definitions can be identified using Lemma 1 and Theorem 1 in Lu et al. (2023). We do not consider the causal estimands $\text{postNDE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140)$ and $\text{postNIE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140)$, because there is no literature to support the identifiability results for these two causal quantities after binarization.

Given the observed evidence $\{X = (1, 1, 1, 1, 1), Y > 140\}$, Table 2 presents the results for each potential risk factor X_k with respect to posterior causal estimands. The first row of Table 2 displays the probabilities of necessity $\text{PN}^*(X_k \Rightarrow \mathcal{E})$ computed after binarizing the outcome variable Y , while the second row presents $\text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140)$. The third to fifth rows present the results for postNDEs, PostNIEs, and PostTCEs considered in this paper. We find that the second and fifth rows of the table show very similar results in terms of sign and ordering. Specifically, HD , E , and D (denoted as X_1 , X_2 , and X_4 , respectively) all have non-zero postTCE values, indicating that they are risk factors for blood pressure. Among them, HD (i.e., X_4) has the largest postTCE, indicating that HD is the most important risk factor for blood pressure. In addition, the value of $\text{PN}^*(X_4 \Rightarrow \mathcal{E})$ is also the largest, further

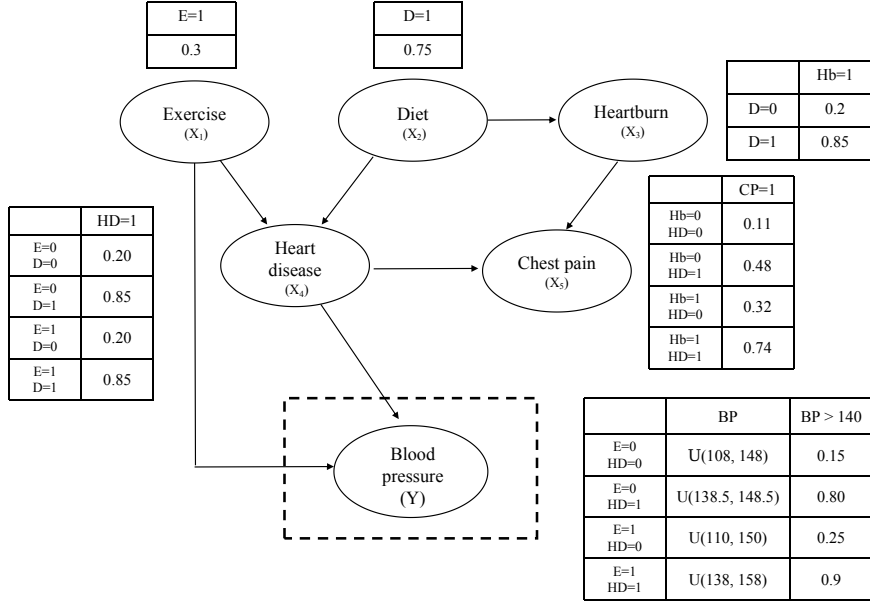


Figure 1: A causal network representing hypertension and its risk factors, where $U(a, b)$ denotes a uniform distribution on the interval $[a, b]$.

confirming the importance of HD as a risk factor. Notably, for HD (i.e., X_4), it can be observed that the postNDE is equal to the postTCE of BP. On the other hand, the postNDE for E (i.e., X_1) is smaller than the postTCE for BP, implying that E has direct and indirect causal effects on BP. The third to fifth rows of the table present the postNDE, postNIE, and postTCE values for Hb and CP (denoted as X_3 and X_5 , respectively), which are all equal to zero. This indicates that they are not risk factors for blood pressure. From the causal network in Figure 1, it can be observed that there is no causal path from Hb (i.e., X_3) and CP (i.e., X_4) to BP. However, $PN^*(X_5 \Rightarrow \mathcal{E})$ is greater than $PN^*(X_1 \Rightarrow \mathcal{E})$ and $PN^*(X_2 \Rightarrow \mathcal{E})$ as shown in the first row.

We also present additional analysis conclusions under different evidence in Section S7 of the supplementary material. To assess the stability of the proposed estimation procedure in Section 4 of the supplementary material, we conduct simulation studies by generating data according to the causal network depicted in Figure 1. The estimated results of Table 2 under different sample sizes were provided. The simulation results indicate negligible biases and small standard errors, particularly for large sample sizes. For detailed information on data generation procedures and simulation results, please refer to Section S8 of the supplementary material. In Section S9 of the supplementary material, we also apply the proposed method to a real dataset (NTP, 2023).

6. Discussion

Dawid et al. (2014) pointed out that statistical inference

about the causes of effects is particularly challenging and is difficult to justify even in randomized experiments. While some prior research has addressed attribution analysis problems with binary outcomes (Pearl, 2000; Dawid et al., 2015; Dawid & Musio, 2022; Li et al., 2023), evaluating the causes of continuous outcomes or effects remains underexplored. The identifiability of ITEs and postICEs with multiple potentially correlated causes is motivated by the identification of ITEs in the case of a single cause (Heckman et al., 1997). However, even if ITEs are identified, it is insufficient for retrospective assessment of a specific cause of effects, as the monotonicity assumption 3.2 is still necessary. In practice, for estimation purposes, we can begin by recovering the counterfactual mappings between different potential outcomes as well as ITEs through quantile regression (Koenker & Bassett, 1978; Heckman et al., 1997; Feng et al., 2020); then, we can utilize the identification expressions in Lemma 3.4 and Theorem 3.6 for nonparametric estimation.

There are several potential directions for future research. Firstly, in addition to continuous outcomes, the attribution analysis of continuous causes is common in practice and is an issue of great interest. Second, the problem of retrospective analysis when multiple causes and multiple continuous outcomes are involved in many medical diagnoses is also worth exploring (Li et al., 2023). Finally, it is also interesting to discuss the bounds of the proposed estimands or to consider sensitivity analysis when the monotonicity assumption does not hold (Tian & Pearl, 2000; Dawid et al., 2024). However, these issues are beyond the scope of this paper and are left for future research.

Impact Statement

This paper proposes a method for causal attribution analysis in the context of continuous outcome variables, which has the potential to improve decision-making in fields such as medicine and policy analysis. By offering a more accurate framework for identifying and estimating posterior intervention effects, our work may lead to more reliable conclusions in various practical scenarios. Ethical considerations include ensuring the robustness and fairness of causal conclusions, especially in domains affecting public health and social well-being. We believe that the methods developed here could contribute to a deeper understanding of complex causal relationships in real-world data.

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Supplementary Material

The supplementary material contains proofs of all theoretical results, identifiability results under a causal network, simulation details of the proposed procedure, and a real data analysis.

S1. The proof of Lemma 3.4

Proof. For any $x \neq x'$, we recall that the definitions of two marginal distributions $F_x(y)$ and $F_{x'}(y)$ are $F_x(y) = \text{pr}(Y \leq y \mid X = x)$ and $F_{x'}(y) = \text{pr}(Y \leq y \mid X = x')$. Let $F_{\epsilon_{p+1}}(t) = \text{pr}(\epsilon_{p+1} \leq t)$ denote the marginal distribution of the error term ϵ_{p+1} . Let $\mathcal{S}_{\epsilon_{p+1}}$ be the support of the error term ϵ_{p+1} . Since the function $m_{p+1}(\cdot, \epsilon_{p+1})$ is strictly monotonic in ϵ_{p+1} under Assumption 3.3, for any $\tau \in \mathcal{S}_{\epsilon_{p+1}}^\circ$ (the interior of $\mathcal{S}_{\epsilon_{p+1}}$), we have,

$$F_{\epsilon_{p+1}}(\tau) = F_{x'}\{m_{p+1}(x', \tau)\} = F_x\{m_{p+1}(x, \tau)\}.$$

Hence, we have $m_{p+1}(x', \tau) \in \mathcal{S}_{Y_{x'}}^\circ$. By the fact that the marginal distribution $F_{x'}(\cdot)$ is continuous and strictly increasing at $m_{p+1}(x', \tau)$, we have,

$$m_{p+1}(x', \tau) = F_{x'}^{-1}[F_x\{m_{p+1}(x, \tau)\}].$$

Let $y = m_{p+1}(x, \tau) \in \mathcal{S}_{Y_x}^\circ$, and we have $\tau = m_{p+1}^{-1}(x, y)$. Then the above equation becomes

$$m_{p+1}(x', \tau) = F_{x'}^{-1}\{F_x(y)\},$$

which shows that $\phi_{x \rightarrow x'}(y)$ is identified on $\mathcal{S}_{Y_x}^\circ$ (the interior of \mathcal{S}_{Y_x}) by

$$\phi_{x \rightarrow x'}(y) = m_{p+1}(x', \tau) = F_{x'}^{-1}\{F_x(y)\}.$$

The function $\phi_{x \rightarrow x'}(y)$ is identified on \mathcal{S}_{Y_x} by its continuous extension. The remaining results follow immediately. \square

S2. The proof of Lemma 3.5

Proof. Without loss of generality, let $x_{k+1:k} = x_{k+1:k}^* = \emptyset$. Let $X_{j:l} = (X_j, \dots, X_l)$ and $x_{j:l} = (x_j, \dots, x_l)$. We first simplify the expression of the conditional probability as follows:

$$\begin{aligned} & \text{pr}\{D_k(a_k, x_k^*) = d_k^* \mid x\} \\ &= \text{pr}\{D_k(a_k, x_k^*) = d_k^* \mid D_k(a_k, x_k) = d_k\} \\ &= \frac{\text{pr}\{D_k(a_k, x_k^*) = d_k^*, D_k(a_k, x_k) = d_k\}}{\text{pr}\{D_k(a_k, x_k) = d_k\}} \\ &= \frac{\text{pr}\{X_{k+1}(a_k, x_k^*) = x_{k+1}^*, \dots, X_p(a_{p-1}, x_{k:p-1}^*) = x_p^*, X_{k+1}(a_k, x_k) = x_{k+1}, \dots, X_p(a_{p-1}, x_{k:p-1}) = x_p\}}{\text{pr}\{X_{k+1}(a_k, x_k) = x_{k+1}, \dots, X_p(a_{p-1}, x_{k:p-1}) = x_p\}} \\ &= \frac{\text{pr}\{X_{k+1}(a_k, x_k^*) = x_{k+1}^*, X_{k+1}(a_k, x_k) = x_{k+1}\} \times \dots \times \text{pr}\{X_p(a_{p-1}, x_{k:p-1}^*) = x_p^*, X_p(a_{p-1}, x_{k:p-1}) = x_p\}}{\text{pr}\{X_{k+1}(a_k, x_k) = x_{k+1}\} \times \dots \times \text{pr}\{X_p(a_{p-1}, x_{k:p-1}) = x_p\}} \\ &= \prod_{s=k+1}^p \text{pr}\{X_s(a_k, x_{k:s-1}^*) = x_s^* \mid X_s(a_k, x_{k:s-1}) = x_s\}. \end{aligned}$$

where the second-to-last equation holds due to Assumption 3.1.

We first consider the case (i), namely $(x_k^*, x_k) = (0, 1)$ and $d_k^* = (x_{k+1}^*, \dots, x_p^*) \preceq d_k = (x_{k+1}, \dots, x_p)$. Under the

monotonicity assumption 3.2, we have,

$$\begin{aligned}
 \Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 1 \mid X_s(a_k, 1, x_{k+1:s-1}) = 1\} &= \frac{\Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 1, X_s(a_k, 1, x_{k+1:s-1}) = 1\}}{\Pr\{X_s(a_k, 1, x_{k+1:s-1}) = 1\}} \\
 &= \frac{\Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 1\}}{\Pr\{X_s(a_k, 1, x_{k+1:s-1}) = 1\}} \\
 &= \frac{\Pr(X_s = 1 \mid a_k, 0, x_{k+1:s-1}^*)}{\Pr(X_s = 1 \mid a_k, 1, x_{k+1:s-1})}. \\
 \Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 0 \mid X_s(a_k, 1, x_{k+1:s-1}) = 1\} &= 1 - \Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 1 \mid X_s(a_k, 1, x_{k+1:s-1}) = 1\}. \\
 \Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 0 \mid X_s(a_k, 1, x_{k+1:s-1}) = 0\} &= \frac{\Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 0, X_s(a_k, 1, x_{k+1:s-1}) = 0\}}{\Pr\{X_s(a_k, 1, x_{k+1:s-1}) = 0\}} \\
 &= \frac{\Pr\{X_s(a_k, 1, x_{k+1:s-1}) = 0\}}{\Pr\{X_s(a_k, 1, x_{k+1:s-1}) = 0\}} \\
 &= 1. \\
 \Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 1 \mid X_s(a_k, 1, x_{k+1:s-1}) = 0\} &= 1 - \Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = 0 \mid X_s(a_k, 1, x_{k+1:s-1}) = 0\} \\
 &= 0.
 \end{aligned}$$

Therefore, we have,

$$\Pr\{X_s(a_k, 0, x_{k+1:s-1}^*) = x_s^* \mid X_s(a_k, 1, x_{k+1:s-1}) = x_s\} = (1 - x_s^*) + (2x_s^* - 1)x_s R_{0s},$$

where

$$R_{0s} = \frac{\Pr(X_s = 1 \mid a_k, 0, x_{k+1}^*, \dots, x_{s-1}^*)}{\Pr(X_s = 1 \mid a_k, 1, x_{k+1}, \dots, x_{s-1})}.$$

We next consider the case (ii), namely $(x_k^*, x_k) = (1, 0)$ and $d_k = (x_{k+1}, \dots, x_p) \preceq d_k^* = (x_{k+1}^*, \dots, x_p^*)$. Under the monotonicity assumption 3.2, we have,

$$\begin{aligned}
 \Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 0 \mid X_s(a_k, 0, x_{k+1:s-1}) = 0\} &= \frac{\Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 0, X_s(a_k, 0, x_{k+1:s-1}) = 0\}}{\Pr\{X_s(a_k, 0, x_{k+1:s-1}) = 0\}} \\
 &= \frac{\Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 0\}}{\Pr\{X_s(a_k, 0, x_{k+1:s-1}) = 0\}} \\
 &= \frac{\Pr(X_s = 0 \mid a_k, 1, x_{k+1:s-1}^*)}{\Pr(X_s = 0 \mid a_k, 0, x_{k+1:s-1})}. \\
 \Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 1 \mid X_s(a_k, 0, x_{k+1:s-1}) = 0\} &= 1 - \Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 0 \mid X_s(a_k, 0, x_{k+1:s-1}) = 0\}. \\
 \Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 1 \mid X_s(a_k, 0, x_{k+1:s-1}) = 1\} &= \frac{\Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 1, X_s(a_k, 0, x_{k+1:s-1}) = 1\}}{\Pr\{X_s(a_k, 0, x_{k+1:s-1}) = 1\}} \\
 &= \frac{\Pr\{X_s(a_k, 0, x_{k+1:s-1}) = 1\}}{\Pr\{X_s(a_k, 0, x_{k+1:s-1}) = 1\}} \\
 &= 1. \\
 \Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 0 \mid X_s(a_k, 0, x_{k+1:s-1}) = 1\} &= 1 - \Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = 1 \mid X_s(a_k, 0, x_{k+1:s-1}) = 1\} \\
 &= 0.
 \end{aligned}$$

Therefore, we have,

$$\Pr\{X_s(a_k, 1, x_{k+1:s-1}^*) = x_s^* \mid X_s(a_k, 0, x_{k+1:s-1}) = x_s\} = x_s^* + (1 - 2x_s^*)(1 - x_s)R_{1s},$$

where

$$R_{1s} = \frac{\Pr(X_s = 0 \mid a_k, 1, x_{k+1}^*, \dots, x_{s-1}^*)}{\Pr(X_s = 0 \mid a_k, 0, x_{k+1}, \dots, x_{s-1})}.$$

□

S3. The proof of Theorem 3.6

Proof. When $x_k = 1$, we first consider $E\{Y_{a_k, x_k^*, D_k(a_k, 1)}\}$:

$$\begin{aligned} E\{Y_{a_k, x_k^*, D_k(a_k, 1)} \mid a_k, 1, d_k, Y \in \mathcal{E}\} &= E\{Y_{a_k, x_k^*, D_k(a_k, 1)} \mid a_k, 1, D_k(a_k, 1) = d_k, Y \in \mathcal{E}\} \\ &= E\{Y_{a_k, x_k^*, d_k} \mid a_k, 1, D_k(a_k, 1) = d_k, Y \in \mathcal{E}\} \\ &= E(Y_{a_k, x_k^*, d_k} \mid a_k, 1, d_k, Y \in \mathcal{E}). \end{aligned}$$

We next consider $E\{Y_{a_k, x_k^*, D_k(a_k, 0)}\}$:

$$\begin{aligned} E\{Y_{a_k, x_k^*, D_k(a_k, 0)} \mid a_k, 1, d_k, Y \in \mathcal{E}\} &= \sum_{d_k^* \preceq d_k} E\{Y_{a_k, x_k^*, d_k^*} \mid a_k, 1, d_k, D_k(a_k, 0) = d_k^*, Y \in \mathcal{E}\} \text{pr}\{D_k(a_k, 0) = d_k^* \mid a_k, 1, d_k\} \\ &= \sum_{d_k^* \preceq d_k} E(Y_{a_k, x_k^*, d_k^*} \mid a_k, 1, d_k, Y \in \mathcal{E}) \text{pr}\{D_k(a_k, 0) = d_k^* \mid a_k, 1, d_k\}, \end{aligned}$$

where the second equality holds due to Assumption 3.1 (ii) and Assumption 3.2. The proofs for $x_k = 0$ follow a similar logic; for simplicity, we omit them. \square

S4. The proof of Corollary 3.7

Proof. Note that because $m_{p+1}(X, \cdot)$ is strictly monotone in ϵ_{p+1} , for any $\tau \in \mathcal{S}_{\epsilon_{p+1}}^\circ$ (the interior of \mathcal{S}_ϵ) and for any $\text{pa}_y^* \neq \text{pa}'_y$, we have:

$$F_{\epsilon_{p+1}}(\tau) = F_{\text{pa}_y^*}\{m_{p+1}(\text{pa}_y^*, \tau)\} = F_{\text{pa}'_y}\{m_{p+1}(\text{pa}'_y, \tau)\}.$$

The remaining proof follows a similar logic as Section S1; we omit it for simplicity. \square

S5. The proof of Corollary 3.8

Proof. Given the Bayesian network, the distribution of each node depends only on its parent nodes. The proof is straightforward, and we omit it for brevity. \square

S6. The proof of Lemma 4.2

Proof. First, we differentiate $\rho_{x \rightarrow x'}(y_{x'}; y_x)$ with respect to y_x . Noting that

$$\begin{aligned} \frac{\partial E(|Y - y| \mid X = x)t}{\partial t} &= -E\{\text{sign}(Y - y) \mid X = x\} = 1 - 2\text{pr}(Y < y \mid X = x) = 1 - 2F_x(y), \\ \frac{\partial E(|Y - t| \mid X = x')}{\partial t} &= -E\{\text{sign}(Y - t) \mid X = x'\} = 1 - 2\text{pr}(Y < t \mid X = x') = 1 - 2F_{x'}(t), \end{aligned}$$

where $\text{sign}(u) \equiv 2 \times \mathbb{I}(u > 0) - 1$. It follows that

$$\frac{\partial \rho_{x \rightarrow x'}(t; y)}{\partial t} = 2 \{F_{x'}(t) - F_x(y)\} = 0.$$

Fix $y_x \in \mathbb{R}$. Note that the marginal distribution $F_{x'}(\cdot)$ is weakly increasing on \mathbb{R} and strictly increasing on $\mathcal{S}_{Y_{x'}}^\circ$. Therefore, $\rho_{x \rightarrow x'}(\cdot; y_x)$ is weakly and strictly convex on \mathbb{R} and $\mathcal{S}_{Y_{x'}}^\circ$. Furthermore, under Assumption 3.3, we know that $\phi_{x \rightarrow x'}(y_x) = F_{x'}^{-1}\{F_x(y_x)\}$. For $y_{x'} \in \mathcal{S}_{Y_{x'}}^\circ$, we have $F_x(y_x) = F_{x'}(y_{x'})$ if and only if $y_{x'} = \phi_{x \rightarrow x'}(y_x)$. Thus, $y_{x'} = \phi_{x \rightarrow x'}(y_x)$ uniquely solves the first-order condition $\frac{\partial}{\partial t} \rho_{x \rightarrow x'}(t; y_x) = 0$. \square

S7. Additional analysis for hypertension example in Section 5

In this section, we present additional analysis conclusions under different evidence. Given the observed evidence $\{X = (1, 1, 1, 0, 1), Y > 140\}$, Table S3 presents the results for the posterior causal estimands with respect to each possible

risk factor X_k . The first two rows of results are directly obtained from Lu et al. (2023) for comparison purposes. In the second row, we observe that $\text{postTCE}^*(X_1 \Rightarrow \mathcal{E} \mid \cdot) = 0.2$, and $\text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid \cdot) = 0$ for $k = 2, \dots, 5$. Our results regarding postTCE are presented in the fifth row, showing a significant increase in blood pressure by 10.483 units when suffering from heart disease, i.e., $\text{postTCE}(X_4 \Rightarrow Y \mid \cdot) = 10.483$. Our results differ from Lu et al. (2023) with respect to X_4 . This is because Lu et al. (2023) additionally requires the monotonicity assumption of the outcome variable for identifiability in the binary outcome case. This assumption ensures that given the evidence $X_4 = 0$ (indicating no heart disease), X_1 is the unique risk factor. However, the perfect rank assumption 3.3 introduced for the continuous variable does not guarantee this. It can also be observed that the probability of necessity is not zero for X_2 , X_3 , and X_5 ; $\text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid \cdot)$, $\text{postTCE}(X_k \Rightarrow Y \mid \cdot)$, and $\text{postNDE}(X_k \Rightarrow Y \mid \cdot)$ are zero when $X_4 = 0$ for $k = 1, 2, 5$. Additionally, it can be observed from the fourth row that $\text{postNIE}(X_k \Rightarrow Y \mid \cdot) = 0$ for all $k = 1, \dots, 5$, suggesting that all causes have no indirect effect on BP. From the causal network in Figure 1, since the evidence indicates no heart disease, i.e., $X_4 = 0$, it can be intuitively understood that X_4 blocks other nodes from transmitting effects to the outcome along the paths.

Table S3: Results of marginal probabilities of necessity and posterior causal estimands based on the evidence $\{X = (1, 1, 1, 0, 1), Y > 140\}$.

	X_1	X_2	X_3	X_4	X_5
$\text{PN}^*(X_k \Rightarrow \mathcal{E})$	0.347	0.230	0.133	#	0.563
$\text{postTCE}^*(X_k \Rightarrow \mathcal{E} \mid x, Y > 140)$	0.200	0	0	0	0
$\text{postNDE}(X_k \Rightarrow Y \mid x, Y > 140)$	2.000	0	0	10.483	0
$\text{postNIE}(X_k \Rightarrow Y \mid x, Y > 140)$	0	0	0	0	0
$\text{postTCE}(X_k \Rightarrow Y \mid x, Y > 140)$	2.000	0	0	10.483	0

Corresponding quantity is undefined.

For the observed evidence $\{X = (1, 0, 1, 1, 1), Y > 140\}$, where $X_2 = 0$ indicates a healthy diet, the values of postTCE and postNDE are shown in Table S4. Comparing it with Table 2, we see that

$$\text{postTCE}\{X_1 \Rightarrow Y \mid X = (1, 0, 1, 1, 1), Y > 140\} = 12.418 > \text{postTCE}\{X_1 \Rightarrow Y \mid X = (1, 1, 1, 1, 1), Y > 140\} = 10.628,$$

and

$$\text{postTCE}\{X_2 \Rightarrow Y \mid X = (1, 0, 1, 1, 1), Y > 140\} = 0 < \text{postTCE}\{X_2 \Rightarrow Y \mid X = (1, 1, 1, 1, 1), Y > 140\} = 4.561.$$

This is because changing $X_2 = 1$ to $X_2 = 0$ in the evidence increases the possibility that X_1 is the cause and decreases the possibility that X_2 is the cause. Similar conclusions are also found in Lu et al. (2023) in the binary case.

S8. Simulation studies for hypertension example in Section 5

S8.1. Data generating details

In this section, we provide additional simulation studies related to Section 5. To generate simulation data that satisfies Figure 1 in Lu et al. (2023), we consider the following data generation process.

- (a) X_1 is Bernoulli with $\text{pr}(X_1 = 1) = 0.3$.
- (b) X_2 is Bernoulli with $\text{pr}(X_2 = 1) = 0.75$.
- (c) Let $G_3 = \{X_3(X_2 = 1), X_3(X_2 = 0)\}$, then G_3 can take values on 00, 01, and 11 under monotonicity assumption 3.2,

$$\text{pr}(G_3 = 00) = 0.15, \text{pr}(G_3 = 01) = 0.65, \text{pr}(G_3 = 11) = 0.2.$$

Here, X_3 is generated as follows:

$$\begin{aligned} X_3 &= 0 \text{ if } G_3 = 00, \\ X_3 &= 1 \text{ if } G_3 = 11, \\ X_3 &= 0 \text{ if } G_3 = 01, X_2 = 0, \\ X_3 &= 0 \text{ if } G_3 = 01, X_2 = 1. \end{aligned}$$

Table S4: Results of marginal probabilities of necessity and posterior causal estimands based on the evidence $\{X = (1, 0, 1, 1, 1), Y > 140\}$.

	X_1	X_2	X_3	X_4	X_5
postTCE* ($X_k \Rightarrow \mathcal{E} \mid x, Y > 140$)	0.449	0	0	0.722	0
postNDE ($X_k \Rightarrow Y \mid x, Y > 140$)	3.424	0	0	16.856	0
postNIE ($X_k \Rightarrow Y \mid x, Y > 140$)	8.994	0	0	0	0
postTCE ($X_k \Rightarrow Y \mid x, Y > 140$)	12.418	0	0	16.856	0

(iv) Let $G_4 = \{X_3(X_1 = 0, X_2 = 0), X_3(X_1 = 0, X_2 = 1), X_3(X_1 = 1, X_2 = 0), X_3(X_1 = 1, X_2 = 1)\}$, then G_4 can take values on 0000, 0001, 0011, 0101, 0111, and 1111 under monotonicity assumption 3.2,

$$\begin{aligned} \text{pr}(G_4 = 0000) &= 0.25, \text{pr}(G_4 = 0001) = 0.10, \text{pr}(G_4 = 0011) = 0.20, \\ \text{pr}(G_4 = 0101) &= 0.10, \text{pr}(G_4 = 0111) = 0.10, \text{pr}(G_4 = 1111) = 0.25. \end{aligned}$$

Here, X_4 is generated as follows:

$$\begin{aligned} X_4 &= 0 \text{ if } G_4 = 0000, \\ X_4 &= 1 \text{ if } G_4 = 0001, X_1 = 1, X_2 = 1, \\ X_4 &= 0 \text{ if } G_4 = 0001, X_1 = 0, X_2 = 1, \\ X_4 &= 0 \text{ if } G_4 = 0001, X_1 = 1, X_2 = 0, \\ X_4 &= 0 \text{ if } G_4 = 0001, X_1 = 0, X_2 = 0, \\ X_4 &= 1 \text{ if } G_4 = 0011, X_1 = 1, \\ X_4 &= 0 \text{ if } G_4 = 0011, X_1 = 0, \\ X_4 &= 1 \text{ if } G_4 = 0101, X_2 = 1, \\ X_4 &= 0 \text{ if } G_4 = 0101, X_2 = 0, \\ X_4 &= 1 \text{ if } G_4 = 0111, X_1 = 1, X_2 = 1, \\ X_4 &= 1 \text{ if } G_4 = 0111, X_1 = 0, X_2 = 1, \\ X_4 &= 1 \text{ if } G_4 = 0111, X_1 = 1, X_2 = 0, \\ X_4 &= 0 \text{ if } G_4 = 0111, X_1 = 0, X_2 = 0, \\ X_4 &= 1 \text{ if } G_4 = 1111. \end{aligned}$$

(d) Let $G_5 = \{X_3(X_3 = 0, X_4 = 0), X_3(X_3 = 0, X_4 = 1), X_3(X_3 = 1, X_4 = 0), X_3(X_3 = 1, X_4 = 1)\}$, then G_5 can take values on 0000, 0001, 0011, 0101, 0111, and 1111 under monotonicity assumption 3.2,

$$\begin{aligned} \text{pr}(G_5 = 0000) &= 0.10, \text{pr}(G_5 = 0001) = 0.05, \text{pr}(G_5 = 0011) = 0.45, \\ \text{pr}(G_5 = 0101) &= 0.05, \text{pr}(G_5 = 0111) = 0.25, \text{pr}(G_5 = 1111) = 0.10. \end{aligned}$$

Here, X_5 is generated as follows:

$$\begin{aligned}
 X_5 &= 0 \text{ if } G_5 = 0000, \\
 X_5 &= 1 \text{ if } G_5 = 0001, X_3 = 1, X_4 = 1, \\
 X_5 &= 0 \text{ if } G_5 = 0001, X_3 = 0, X_4 = 1, \\
 X_5 &= 0 \text{ if } G_5 = 0001, X_3 = 1, X_4 = 0, \\
 X_5 &= 0 \text{ if } G_5 = 0001, X_3 = 0, X_4 = 0, \\
 X_5 &= 1 \text{ if } G_5 = 0011, X_3 = 1, \\
 X_5 &= 0 \text{ if } G_5 = 0011, X_3 = 0, \\
 X_5 &= 1 \text{ if } G_5 = 0101, X_4 = 1, \\
 X_5 &= 0 \text{ if } G_5 = 0101, X_4 = 0, \\
 X_5 &= 1 \text{ if } G_5 = 0111, X_3 = 1, X_4 = 1, \\
 X_5 &= 1 \text{ if } G_5 = 0111, X_3 = 0, X_4 = 1, \\
 X_5 &= 1 \text{ if } G_5 = 0111, X_3 = 1, X_4 = 0, \\
 X_5 &= 0 \text{ if } G_5 = 0111, X_3 = 0, X_4 = 0, \\
 X_5 &= 1 \text{ if } G_5 = 1111.
 \end{aligned}$$

The above data generation process will result in an observation distribution exactly identical to [Lu et al. \(2023\)](#) and will be utilized to compute the true values of posterior causal estimands considered in this paper.

S8.2. The simulation results for posterior causal estimands

In this section, we conducted simulation studies to assess the performance of the proposed procedure in finite samples. We generated numerical examples corresponding to Figure 1 of the main text using the method described in Section S8.1. Using data generated from 2000000 samples, we obtained the estimated results presented in Tables 2, S3, and S4 of the main text, as well as the true values provided in Tables S5 and S6 of the supplementary material. Additionally, we considered sample sizes of $n = 1000, 2000$, and 10000 , and averaged the results over 500 repetitions. These simulation results demonstrate the stability of the proposed identification expressions. Specifically, we observed that the estimated values are close to the true values, and the standard errors are relatively small. As the sample size increases, both the bias and standard errors decrease.

- (i) Table S5 presents the true values and estimated results of the posterior intervention causal effect based on different evidence.
- (ii) Table S6 presents the true values of the posterior natural direct and indirect causal effect based on different evidence.
- (iii) Tables S7-S9 present the estimated results of the posterior direct effect based on different evidence.
- (iv) Tables S10-S12 present the estimated results of the posterior indirect effect based on different evidence.

Table S5: Posterior intervention causal effect based on different pieces of evidence.

	PostICE($Y'_x \mid x, Y > 140$)	$(x_1, x_4) = (0, 0)$	$(x_1, x_4) = (0, 1)$	$(x_1, x_4) = (1, 0)$	$(x_1, x_4) = (1, 1)$
True values	$(x'_1, x'_4) = (0, 0)$	0	-13.26	-2	-19
	$(x'_1, x'_4) = (0, 1)$	3.50	0	2.25	-5
	$(x'_1, x'_4) = (1, 0)$	2	-11.26	0	-17
	$(x'_1, x'_4) = (1, 1)$	12	5.25	10.5	0
$n = 1000$	$(x'_1, x'_4) = (0, 0)$	0.000 (0.000)	-13.252 (0.722)	-1.757 (1.210)	-18.955 (0.769)
	$(x'_1, x'_4) = (0, 1)$	3.477 (0.317)	0.000 (0.000)	2.391 (0.699)	-4.998 (0.432)
	$(x'_1, x'_4) = (1, 0)$	1.561 (1.129)	-11.532 (1.496)	0.000 (0.000)	-17.224 (1.480)
	$(x'_1, x'_4) = (1, 1)$	11.967 (0.435)	5.227 (0.518)	10.676 (0.835)	0.000 (0.000)
$n = 2000$	$(x'_1, x'_4) = (0, 0)$	0.000 (0.000)	-13.224 (0.554)	-1.921 (0.928)	-18.962 (0.602)
	$(x'_1, x'_4) = (0, 1)$	3.463 (0.216)	0.000 (0.000)	2.273 (0.495)	-4.997 (0.300)
	$(x'_1, x'_4) = (1, 0)$	1.808 (0.862)	-11.322 (1.031)	0.000 (0.000)	-17.057 (1.073)
	$(x'_1, x'_4) = (1, 1)$	11.944 (0.296)	5.240 (0.361)	10.518 (0.631)	0.000 (0.000)
$n = 10000$	$(x'_1, x'_4) = (0, 0)$	0.000 (0.0000)	-13.261 (0.2636)	-1.950 (0.3588)	-19.036 (0.2550)
	$(x'_1, x'_4) = (0, 1)$	3.503 (0.0992)	0.000 (0.0000)	2.291 (0.1965)	-5.005 (0.1383)
	$(x'_1, x'_4) = (1, 0)$	1.926 (0.3298)	-11.307 (0.4901)	0.000 (0.0000)	-17.076 (0.4652)
	$(x'_1, x'_4) = (1, 1)$	11.995 (0.1558)	5.261 (0.1576)	10.543 (0.2530)	0.000 (0.0000)

S9. Real data analysis: risk factors for abnormal weight

In this section, we apply the proposed method to a real dataset from the developmental toxicology experiments conducted by the National Toxicology Program (NTP) (NTP, 2023). The primary objective of this study is to analyze whether tris(1-chloro-2-propyl) phosphate (TCPP) is a risk factor for abnormal weight loss in B6C3F1/N mice, or if there are other causes that are the potential risks. In this experiment, a total of 120 mice were randomly exposed to six different dose levels of TCPP via dosed feed: 0, 1250, 2500, 5000, 10000, or 20000 ppm, for a duration of 3 months. Each pup’s data includes gender (male/female), weekly body weights, organ weights, and whether organs exhibit pathology. In our analysis, let X_1 represent the gender of the mice, where $X_1 = 0$ indicates female mice and $X_1 = 1$ indicates male mice. Let X_2 denote the dose, where $X_2 = 0$ represents exposure to the low dose group including 0, 1250, and 2500 ppm, and $X_2 = 1$ represents exposure to the high dose group including 5000, 10000, and 20000 ppm. Let X_3 denote whether the liver or kidney exhibits pathology, where $X_3 = 0$ indicates no pathology and $X_3 = 1$ indicates pathology. We choose the body weight at the end of three months as the outcome Y . In this analysis, we focus on assessing the potential risk factors affecting the body weight of underweight mice, indicated by the event $\mathcal{E} = \mathbb{I}(Y < 27)$.

We first use the R package “bnlearn” to construct a Bayesian network based on the collected data as shown in Figure S2. It is clear that dose X_2 indirectly affects body weight Y through the organ disease X_3 . Since both gender (i.e., X_1) and dose (i.e., X_2) can be considered as randomized trials, we can estimate the causal effect of X_1 on Y using the difference in means estimator, i.e., $E(Y_{X_1=1} - Y_{X_1=0}) \approx 6.09$; which implies that males are heavier. Similarly, we can estimate the causal effect of X_2 on Y to be -3.353, i.e., $E(Y_{X_2=1} - Y_{X_2=0}) \approx -3.353$, which suggests that the higher dose lead to weight loss. Figure S2 also provides a simple descriptive statistical analysis of these potential risk factors. We observe that as the toxin level increases, the occurrence rate of organ disease also increases. In addition, males showed higher variability in organ abnormalities compared to females, as demonstrated in previous studies (Bianco et al., 2023). These empirical findings align with monotonicity assumption 3.2, i.e., $X_3(0, 0) \leq \{X_3(0, 1), X_3(1, 0)\} \leq X_3(1, 1)$. In this data analysis, we choose not to binarize the outcome and, therefore, do not report the results of the binarized posterior causal effects, as the outcome variable fails to satisfy the monotonicity assumption regarding the causes (Assumption 2(b) in Lu et al. (2023)).

We now present the estimation results of postICEs based on various observed evidence in Table S13. According to Corollary 3.7 and Figure S2, we know that the postICEs are only related to the parent nodes of body weight, namely X_1 (gender) and X_3 (organ disease). Considering the evidence $(X_1, X_3, \mathcal{E}) = (0, 1, Y < 27)$, we observe $\text{PostICE}(Y_{x'} \mid x, Y < 27) = 10.51$, which indicates that changing from male mice without organ disease (i.e., $(x'_1, x'_3) = (1, 0)$) to female mice with organ disease results in the most significant weight loss, totaling 10.51. For the evidence $(x_1, x_3, \mathcal{E}) = (0, 0, Y < 27)$, we observe $\text{PostICE}(Y_{x'} \mid x, Y < 27) = -1.42$, which indicates that changing from female mice with organ disease (i.e., $(x'_1, x'_3) = (0, 1)$) to female mice without organ disease results in a slight decrease of about 1.42. In summary, for a given posterior evidence, changing from female mice ($X_1 = 0$) to male mice ($X_1 = 1$) results in an increase in body weight,

Table S6: True values of posterior natural direct effect and posterior natural indirect effect.

Case	Evidence (x, \mathcal{E})	Posterior natural direct effect					Posterior natural indirect effect				
		X_1	X_2	X_3	X_4	X_5	X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0, \mathcal{E})	2.000	0.000	0.000	3.482	0.000	3.885	0.918	0.000	0.000	0.000
2	(0,0,0,0,1, \mathcal{E})	2.000	0.000	0.000	3.331	0.000	3.694	0.796	0.000	0.000	0.000
3	(0,0,0,1,0, \mathcal{E})	5.197	0.000	0.000	13.408	0.000	0.000	0.000	0.000	0.000	0.000
4	(0,0,0,1,1, \mathcal{E})	5.201	0.000	0.000	13.396	0.000	0.000	0.000	0.000	0.000	0.000
5	(0,0,1,0,0, \mathcal{E})	2.000	0.000	0.000	3.539	0.000	4.460	0.992	0.000	0.000	0.000
6	(0,0,1,0,1, \mathcal{E})	2.000	0.000	0.000	3.434	0.000	3.908	0.838	0.000	0.000	0.000
7	(0,0,1,1,0, \mathcal{E})	5.416	0.000	0.000	12.751	0.000	0.000	0.000	0.000	0.000	0.000
8	(0,0,1,1,1, \mathcal{E})	5.171	0.000	0.000	13.486	0.000	0.000	0.000	0.000	0.000	0.000
9	(0,1,0,0,0, \mathcal{E})	2.000	0.000	0.000	3.455	0.000	5.655	0.000	0.000	0.000	0.000
10	(0,1,0,0,1, \mathcal{E})	2.000	0.000	0.000	3.585	0.000	5.474	0.000	0.000	0.000	0.000
11	(0,1,0,1,0, \mathcal{E})	5.126	0.000	0.000	13.621	0.000	0.000	5.926	0.000	0.000	0.000
12	(0,1,0,1,1, \mathcal{E})	5.291	0.000	0.000	13.127	0.000	0.000	5.950	0.000	0.000	0.000
13	(0,1,1,0,0, \mathcal{E})	2.000	0.000	0.000	3.547	0.000	5.664	0.000	0.000	0.000	0.000
14	(0,1,1,0,1, \mathcal{E})	2.000	0.000	0.000	3.483	0.000	5.407	0.000	0.000	0.000	0.000
15	(0,1,1,1,0, \mathcal{E})	5.192	0.000	0.000	13.425	0.000	0.000	5.874	0.000	0.000	0.000
16	(0,1,1,1,1, \mathcal{E})	5.246	0.000	0.000	13.263	0.000	0.000	5.916	0.000	0.000	0.000
17	(1,0,0,0,0, \mathcal{E})	2.000	0.000	0.000	10.490	0.000	0.000	4.602	0.000	0.000	0.000
18	(1,0,0,0,1, \mathcal{E})	2.000	0.000	0.000	10.365	0.000	0.000	4.864	0.000	0.000	0.000
19	(1,0,0,1,0, \mathcal{E})	3.342	0.000	0.000	16.955	0.000	9.184	0.000	0.000	0.000	0.000
20	(1,0,0,1,1, \mathcal{E})	3.304	0.000	0.000	17.188	0.000	9.477	0.000	0.000	0.000	0.000
21	(1,0,1,0,0, \mathcal{E})	2.000	0.000	0.000	10.569	0.000	0.000	4.793	0.000	0.000	0.000
22	(1,0,1,0,1, \mathcal{E})	2.000	0.000	0.000	10.645	0.000	0.000	3.802	0.000	0.000	0.000
23	(1,0,1,1,0, \mathcal{E})	3.504	0.000	0.000	17.188	0.000	8.130	0.000	0.000	0.000	0.000
24	(1,0,1,1,1, \mathcal{E})	3.424	0.000	0.000	16.856	0.000	8.994	0.000	0.000	0.000	0.000
25	(1,1,0,0,0, \mathcal{E})	2.000	0.000	0.000	10.363	0.000	0.000	0.000	0.000	0.000	0.000
26	(1,1,0,0,1, \mathcal{E})	2.000	0.000	0.000	10.586	0.000	0.000	0.000	0.000	0.000	0.000
27	(1,1,0,1,0, \mathcal{E})	3.867	0.000	0.000	16.809	0.000	6.740	4.306	0.000	0.000	0.000
28	(1,1,0,1,1, \mathcal{E})	3.799	0.000	0.000	17.045	0.000	6.692	4.251	0.000	0.000	0.000
29	(1,1,1,0,0, \mathcal{E})	2.000	0.000	0.000	10.463	0.000	0.000	0.000	0.000	0.000	0.000
30	(1,1,1,0,1, \mathcal{E})	2.000	0.000	0.000	10.483	0.000	0.000	0.000	0.000	0.000	0.000
31	(1,1,1,1,0, \mathcal{E})	3.837	0.000	0.000	16.993	0.000	6.595	4.603	0.000	0.000	0.000
32	(1,1,1,1,1, \mathcal{E})	3.823	0.000	0.000	17.023	0.000	6.805	4.561	0.000	0.000	0.000

while changing from mice without organ disease ($X_3 = 0$) to mice with organ disease ($X_3 = 1$) results in a decrease in body weight in most cases.

Table S14 presents the estimation results of the posterior causal estimands for each potential risk factor. Given the evidence $(X_1, X_2, X_3, \mathcal{E}) = (1, 1, 1, Y < 27)$, we find that the PostNDE value for gender X_1 is the highest, indicating that gender might be the most important direct factor, followed by organ disease X_3 . The PostNDE value for toxin dose X_2 is 0, which is consistent with the conclusions drawn from the Bayesian network in Figure S2. In addition, given the evidence $(X_1, X_2, X_3, \mathcal{E}) = (1, 1, 1, Y < 27)$, we observed that both gender X_1 and toxin dose X_2 have an indirect effect on the outcome, with the absolute value of PostNIE for gender X_1 being significantly greater than that for toxin dose X_2 , while organ disease X_3 has no indirect effect, suggesting that gender may be the most important indirect factor influencing body weight among the three factors, followed by toxin dose X_2 . However, in terms of the postTCE, organ disease X_3 has the largest value, suggesting that organ disease X_3 is the most important total risk factor for low body weight, followed by toxin dose X_2 , while gender X_1 has the least effect.

Under the evidence $(1, 0, 0, Y < 27)$ and $(0, 1, 1, Y < 27)$, toxin dose X_2 exhibits the most significant PostNIE value for weight loss, indicating that toxin dose X_2 is the most important indirect risk factor in these situations. In summary, for most evidence in Table S14, either gender X_1 or organ disease X_3 has the highest absolute PostTCE value, especially X_3 , suggesting that organ disease is the most important risk factor leading to weight loss.

Table S7: The simulation results for the postNDE under a sample size of 1000.

Case	Evidence (x, \mathcal{E})	PostNDE($X_k \Rightarrow Y \mid x, Y > 140$)				
		X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0,0, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
2	(0,0,0,0,1, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
3	(0,0,0,1,0, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
4	(0,0,0,1,1, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
5	(0,0,1,0,0, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
6	(0,0,1,0,1, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
7	(0,0,0,1,0, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
8	(0,0,1,1,1, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
9	(0,1,0,0,0, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
10	(0,1,0,0,1, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
11	(0,1,0,1,0, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
12	(0,1,0,1,1, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
13	(0,1,1,0,0, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
14	(0,1,1,0,1, \mathcal{E})	1.561 (1.134)	0.000 (0.000)	0.000 (0.000)	3.489 (0.318)	0.000 (0.000)
15	(0,1,1,1,0, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
16	(0,1,1,1,1, \mathcal{E})	5.226 (0.519)	0.000 (0.000)	0.000 (0.000)	13.255 (0.722)	0.000 (0.000)
17	(1,0,0,0,0, \mathcal{E})	1.755 (1.194)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
18	(1,0,0,0,1, \mathcal{E})	1.756 (1.197)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
19	(1,0,0,1,0, \mathcal{E})	3.230 (0.903)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)
20	(1,0,0,1,1, \mathcal{E})	3.230 (0.903)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)
21	(1,0,1,0,0, \mathcal{E})	1.755 (1.194)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
22	(1,0,1,0,1, \mathcal{E})	1.753 (1.193)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
23	(1,0,1,1,0, \mathcal{E})	3.371 (1.153)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)
24	(1,0,1,1,1, \mathcal{E})	3.365 (1.148)	0.000 (0.000)	0.000 (0.000)	17.217 (1.478)	0.000 (0.000)
25	(1,1,0,0,0, \mathcal{E})	1.755 (1.194)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
26	(1,1,0,0,1, \mathcal{E})	1.804 (1.124)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
27	(1,1,0,1,0, \mathcal{E})	3.732 (0.740)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)
28	(1,1,0,1,1, \mathcal{E})	3.732 (0.740)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)
29	(1,1,1,0,0, \mathcal{E})	1.755 (1.194)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
30	(1,1,1,0,1, \mathcal{E})	1.755 (1.194)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
31	(1,1,1,1,0, \mathcal{E})	3.692 (0.667)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)
32	(1,1,1,1,1, \mathcal{E})	3.692 (0.667)	0.000 (0.000)	0.000 (0.000)	17.220 (1.479)	0.000 (0.000)

Table S8: The simulation results for the postNDE under a sample size of 2000.

Case	Evidence (x, \mathcal{E})	PostNDE($X_k \Rightarrow Y \mid x, Y > 140$)				
		X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0,0, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
2	(0,0,0,0,1, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
3	(0,0,0,1,0, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
4	(0,0,0,1,1, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
5	(0,0,1,0,0, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.914 (0.200)	3.469 (0.217)	0.000 (0.000)
6	(0,0,1,0,1, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
7	(0,0,1,1,0, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
8	(0,0,1,1,1, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
9	(0,1,0,0,0, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
10	(0,1,0,0,1, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
11	(0,1,0,1,0, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
12	(0,1,0,1,1, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
13	(0,1,1,0,0, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
14	(0,1,1,0,1, \mathcal{E})	1.808 (0.863)	0.000 (0.000)	0.000 (0.000)	3.469 (0.217)	0.000 (0.000)
15	(0,1,1,1,0, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
16	(0,1,1,1,1, \mathcal{E})	5.239 (0.361)	0.000 (0.000)	0.000 (0.000)	13.226 (0.554)	0.000 (0.000)
17	(1,0,0,0,0, \mathcal{E})	1.920 (0.921)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
18	(1,0,0,0,1, \mathcal{E})	1.922 (0.922)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
19	(1,0,0,1,0, \mathcal{E})	3.310 (0.622)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
20	(1,0,0,1,1, \mathcal{E})	3.310 (0.622)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
21	(1,0,1,0,0, \mathcal{E})	1.920 (0.921)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
22	(1,0,1,0,1, \mathcal{E})	1.920 (0.921)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
23	(1,0,1,1,0, \mathcal{E})	3.318 (0.724)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
24	(1,0,1,1,1, \mathcal{E})	3.318 (0.724)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
25	(1,1,0,0,0, \mathcal{E})	1.920 (0.921)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
26	(1,1,0,0,1, \mathcal{E})	1.804 (1.124)	0.000 (0.000)	0.000 (0.000)	10.641 (0.821)	0.000 (0.000)
27	(1,1,0,1,0, \mathcal{E})	3.753 (0.523)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
28	(1,1,0,1,1, \mathcal{E})	3.753 (0.523)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
29	(1,1,1,0,0, \mathcal{E})	1.920 (0.921)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
30	(1,1,1,0,1, \mathcal{E})	1.920 (0.921)	0.000 (0.000)	0.000 (0.000)	10.501 (0.625)	0.000 (0.000)
31	(1,1,1,1,0, \mathcal{E})	3.760 (0.461)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)
32	(1,1,1,1,1, \mathcal{E})	3.760 (0.461)	0.000 (0.000)	0.000 (0.000)	17.055 (1.073)	0.000 (0.000)

Table S9: The simulation results for the postNDE under a sample size of 10000.

Case	Evidence (x, \mathcal{E})	postNDE($X_k \Rightarrow Y \mid x, Y > 140$)				
		X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0,0, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
2	(0,0,0,0,1, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
3	(0,0,0,1,0, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
4	(0,0,0,1,1, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
5	(0,0,1,0,0, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
6	(0,0,1,0,1, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
7	(0,0,0,1,0, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
8	(0,0,1,1,1, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
9	(0,1,0,0,0, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
10	(0,1,0,0,1, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
11	(0,1,0,1,0, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
12	(0,1,0,1,1, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
13	(0,1,1,0,0, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
14	(0,1,1,0,1, \mathcal{E})	1.961(0.351)	0.000(0.000)	0.000(0.000)	3.504(0.102)	0.000(0.000)
15	(0,1,1,1,0, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
16	(0,1,1,1,1, \mathcal{E})	5.259(0.171)	0.000(0.000)	0.000(0.000)	13.259(0.261)	0.000(0.000)
17	(1,0,0,0,0, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
18	(1,0,0,0,1, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
19	(1,0,0,1,0, \mathcal{E})	3.357(0.267)	0.000(0.000)	0.000(0.000)	17.050(0.466)	0.000(0.000)
20	(1,0,0,1,1, \mathcal{E})	3.357(0.267)	0.000(0.000)	0.000(0.000)	17.050(0.466)	0.000(0.000)
21	(1,0,1,0,0, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
22	(1,0,1,0,1, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
23	(1,0,1,1,0, \mathcal{E})	3.364(0.322)	0.000(0.000)	0.000(0.000)	17.050(0.466)	0.000(0.000)
24	(1,0,1,1,1, \mathcal{E})	3.364(0.322)	0.000(0.000)	0.000(0.000)	17.050(0.466)	0.000(0.000)
25	(1,1,0,0,0, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
26	(1,1,0,0,1, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
27	(1,1,0,1,0, \mathcal{E})	3.796(0.229)	0.000(0.000)	0.000(0.000)	17.050(0.466)	0.000(0.000)
28	(1,1,0,1,1, \mathcal{E})	3.798(0.228)	0.000(0.000)	0.000(0.000)	17.045(0.466)	0.000(0.000)
29	(1,1,1,0,0, \mathcal{E})	1.983(0.382)	0.000(0.000)	0.000(0.000)	10.513(0.267)	0.000(0.000)
30	(1,1,1,0,1, \mathcal{E})	1.978(0.381)	0.000(0.000)	0.000(0.000)	10.519(0.269)	0.000(0.000)
31	(1,1,1,1,0, \mathcal{E})	3.798(0.228)	0.000(0.000)	0.000(0.000)	17.047(0.470)	0.000(0.000)
32	(1,1,1,1,1, \mathcal{E})	3.790(0.195)	0.000(0.000)	0.000(0.000)	17.076(0.465)	0.000(0.000)

Table S10: The simulation results for the postNIE under a sample size of 1000.

Case	Evidence (x, \mathcal{E})	PostNIE($X_k \Rightarrow Y \mid x, Y > 140$)				
		X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0,0, \mathcal{E})	4.157 (1.094)	0.926 (0.186)	-0.001 (0.376)	0.000 (0.000)	0.000 (0.000)
2	(0,0,0,0,1, \mathcal{E})	4.159 (1.093)	0.926 (0.186)	-0.001 (0.376)	0.000 (0.000)	0.000 (0.000)
3	(0,0,0,1,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
4	(0,0,0,1,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
5	(0,0,1,0,0, \mathcal{E})	4.058 (1.995)	0.903 (0.286)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
6	(0,0,1,0,1, \mathcal{E})	4.082 (2.013)	0.901 (0.286)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
7	(0,0,0,1,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-1.186 (5.424)	0.000 (0.000)	0.000 (0.000)
8	(0,0,1,1,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-1.186 (5.424)	0.000 (0.000)	0.000 (0.000)
9	(0,1,0,0,0, \mathcal{E})	5.580 (1.742)	0.000 (0.000)	-0.056 (0.429)	0.000 (0.000)	0.000 (0.000)
10	(0,1,0,0,1, \mathcal{E})	5.580 (1.742)	0.000 (0.000)	-0.056 (0.429)	0.000 (0.000)	0.000 (0.000)
11	(0,1,0,1,0, \mathcal{E})	0.000 (0.000)	5.801 (1.560)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
12	(0,1,0,1,1, \mathcal{E})	0.000 (0.000)	5.801 (1.560)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
13	(0,1,1,0,0, \mathcal{E})	5.707 (0.874)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
14	(0,1,1,0,1, \mathcal{E})	5.707 (0.874)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
15	(0,1,1,1,0, \mathcal{E})	0.000 (0.000)	5.870 (1.056)	-0.094 (1.876)	0.000 (0.000)	0.000 (0.000)
16	(0,1,1,1,1, \mathcal{E})	0.000 (0.000)	5.870 (1.056)	-0.094 (1.876)	0.000 (0.000)	0.000 (0.000)
17	(1,0,0,0,0, \mathcal{E})	0.000 (0.000)	4.616 (1.206)	-0.266 (3.567)	0.000 (0.000)	0.000 (0.000)
18	(1,0,0,0,1, \mathcal{E})	0.000 (0.000)	4.616 (1.206)	-0.246 (3.591)	0.000 (0.000)	0.000 (0.000)
19	(1,0,0,1,0, \mathcal{E})	9.322 (1.777)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
20	(1,0,0,1,1, \mathcal{E})	9.322 (1.777)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
21	(1,0,1,0,0, \mathcal{E})	0.000 (0.000)	4.544 (2.096)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
22	(1,0,1,0,1, \mathcal{E})	0.000 (0.000)	4.159 (2.725)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
23	(1,0,1,1,0, \mathcal{E})	8.688 (4.183)	0.000 (0.000)	-1.352 (6.608)	0.000 (0.000)	0.000 (0.000)
24	(1,0,1,1,1, \mathcal{E})	8.708 (4.163)	0.000 (0.000)	-1.345 (6.613)	0.000 (0.000)	0.000 (0.000)
25	(1,1,0,0,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-1.177 (5.164)	0.000 (0.000)	0.000 (0.000)
26	(1,1,0,0,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-1.177 (5.164)	0.000 (0.000)	0.000 (0.000)
27	(1,1,0,1,0, \mathcal{E})	6.711 (1.876)	4.408 (2.022)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
28	(1,1,0,1,1, \mathcal{E})	6.711 (1.876)	4.408 (2.022)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
29	(1,1,1,0,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
30	(1,1,1,0,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
31	(1,1,1,1,0, \mathcal{E})	6.902 (0.902)	4.582 (1.508)	0.043 (1.996)	0.000 (0.000)	0.000 (0.000)
32	(1,1,1,1,1, \mathcal{E})	6.902 (0.902)	4.588 (1.503)	0.043 (1.996)	0.000 (0.000)	0.000 (0.000)

Table S11: The simulation results for the postNIE under a sample size of 2000.

Case	Evidence (x, \mathcal{E})	PostNIE($X_k \Rightarrow Y \mid x, Y > 140$)				
		X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0, \mathcal{E})	4.059 (0.748)	0.921 (0.120)	-0.004 (0.256)	0.000 (0.000)	0.000 (0.000)
2	(0,0,0,0,1, \mathcal{E})	4.059 (0.748)	0.921 (0.120)	-0.004 (0.256)	0.000 (0.000)	0.000 (0.000)
3	(0,0,0,1,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
4	(0,0,0,1,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
5	(0,0,1,0,0, \mathcal{E})	4.120 (1.418)	0.914 (0.200)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
6	(0,0,1,0,1, \mathcal{E})	4.120 (1.418)	0.914 (0.200)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
7	(0,0,0,1,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.576 (3.295)	0.000 (0.000)	0.000 (0.000)
8	(0,0,1,1,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.576 (3.295)	0.000 (0.000)	0.000 (0.000)
9	(0,1,0,0,0, \mathcal{E})	5.531 (1.246)	0.000 (0.000)	-0.005 (0.271)	0.000 (0.000)	0.000 (0.000)
10	(0,1,0,0,1, \mathcal{E})	5.531 (1.246)	0.000 (0.000)	-0.005 (0.271)	0.000 (0.000)	0.000 (0.000)
11	(0,1,0,1,0, \mathcal{E})	0.000 (0.000)	5.808 (1.006)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
12	(0,1,0,1,1, \mathcal{E})	0.000 (0.000)	5.808 (1.006)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
13	(0,1,1,0,0, \mathcal{E})	5.544 (0.653)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
14	(0,1,1,0,1, \mathcal{E})	5.544 (0.653)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
15	(0,1,1,1,0, \mathcal{E})	0.000 (0.000)	5.887 (0.746)	0.041 (1.254)	0.000 (0.000)	0.000 (0.000)
16	(0,1,1,1,1, \mathcal{E})	0.000 (0.000)	5.887 (0.746)	0.041 (1.254)	0.000 (0.000)	0.000 (0.000)
17	(1,0,0,0,0, \mathcal{E})	0.000 (0.000)	4.606 (0.828)	0.025 (2.427)	0.000 (0.000)	0.000 (0.000)
18	(1,0,0,0,1, \mathcal{E})	0.000 (0.000)	4.606 (0.828)	0.025 (2.427)	0.000 (0.000)	0.000 (0.000)
19	(1,0,0,1,0, \mathcal{E})	9.308 (1.114)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
20	(1,0,0,1,1, \mathcal{E})	9.308 (1.114)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
21	(1,0,1,0,0, \mathcal{E})	0.000 (0.000)	4.406 (1.524)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
22	(1,0,1,0,1, \mathcal{E})	0.000 (0.000)	4.296 (1.620)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
23	(1,0,1,1,0, \mathcal{E})	9.227 (2.246)	0.000 (0.000)	-0.376 (3.501)	0.000 (0.000)	0.000 (0.000)
24	(1,0,1,1,1, \mathcal{E})	9.227 (2.246)	0.000 (0.000)	-0.376 (3.501)	0.000 (0.000)	0.000 (0.000)
25	(1,1,0,0,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.546 (2.909)	0.000 (0.000)	0.000 (0.000)
26	(1,1,0,0,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.546 (2.909)	0.000 (0.000)	0.000 (0.000)
27	(1,1,0,1,0, \mathcal{E})	6.819 (1.306)	4.437 (1.473)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
28	(1,1,0,1,1, \mathcal{E})	6.819 (1.306)	4.437 (1.473)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
29	(1,1,1,0,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
30	(1,1,1,0,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
31	(1,1,1,1,0, \mathcal{E})	6.827 (0.693)	4.489 (1.034)	0.007 (1.395)	0.000 (0.000)	0.000 (0.000)
32	(1,1,1,1,1, \mathcal{E})	6.827 (0.693)	4.489 (1.034)	0.007 (1.395)	0.000 (0.000)	0.000 (0.000)

Table S12: The simulation results for postNIE under sample size 10000.

Case	Evidence (x, \mathcal{E})	PostNIE($X_k \Rightarrow Y \mid x, Y > 140$)				
		X_1	X_2	X_3	X_4	X_5
1	(0,0,0,0, \mathcal{E})	4.004 (0.331)	0.934 (0.057)	0.001 (0.122)	0.000 (0.000)	0.000 (0.000)
2	(0,0,0,0,1, \mathcal{E})	4.004 (0.331)	0.934 (0.057)	0.001 (0.122)	0.000 (0.000)	0.000 (0.000)
3	(0,0,0,1,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
4	(0,0,0,1,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
5	(0,0,1,0,0, \mathcal{E})	3.989 (0.590)	0.931 (0.088)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
6	(0,0,1,0,1, \mathcal{E})	3.989 (0.590)	0.931 (0.088)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
7	(0,0,0,1,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.098 (1.409)	0.000 (0.000)	0.000 (0.000)
8	(0,0,1,1,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.098 (1.409)	0.000 (0.000)	0.000 (0.000)
9	(0,1,0,0,0, \mathcal{E})	5.458 (0.491)	0.000 (0.000)	-0.004 (0.256)	0.000 (0.000)	0.000 (0.000)
10	(0,1,0,0,1, \mathcal{E})	5.458 (0.491)	0.000 (0.000)	-0.004 (0.126)	0.000 (0.000)	0.000 (0.000)
11	(0,1,0,1,0, \mathcal{E})	0.000 (0.000)	5.880 (0.466)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
12	(0,1,0,1,1, \mathcal{E})	0.000 (0.000)	5.880 (0.466)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
13	(0,1,1,0,0, \mathcal{E})	5.472 (0.259)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
14	(0,1,1,0,1, \mathcal{E})	5.472 (0.259)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
15	(0,1,1,1,0, \mathcal{E})	0.000 (0.000)	5.886 (0.336)	-0.005 (0.577)	0.000 (0.000)	0.000 (0.000)
16	(0,1,1,1,1, \mathcal{E})	0.000 (0.000)	5.886 (0.336)	-0.005 (0.577)	0.000 (0.000)	0.000 (0.000)
17	(1,0,0,0,0, \mathcal{E})	0.000 (0.000)	4.664 (0.346)	-0.037 (1.038)	0.000 (0.000)	0.000 (0.000)
18	(1,0,0,0,1, \mathcal{E})	0.000 (0.000)	4.664 (0.346)	-0.037 (1.038)	0.000 (0.000)	0.000 (0.000)
19	(1,0,0,1,0, \mathcal{E})	9.275 (0.501)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
20	(1,0,0,1,1, \mathcal{E})	9.275 (0.501)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
21	(1,0,1,0,0, \mathcal{E})	0.000 (0.000)	4.640 (0.586)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
22	(1,0,1,0,1, \mathcal{E})	0.000 (0.000)	4.640 (0.586)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
23	(1,0,1,1,0, \mathcal{E})	9.220 (1.009)	0.000 (0.000)	-0.114 (1.414)	0.000 (0.000)	0.000 (0.000)
24	(1,0,1,1,1, \mathcal{E})	9.220 (1.009)	0.000 (0.000)	-0.114 (1.414)	0.000 (0.000)	0.000 (0.000)
25	(1,1,0,0,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.069 (1.093)	0.000 (0.000)	0.000 (0.000)
26	(1,1,0,0,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	-0.069 (1.093)	0.000 (0.000)	0.000 (0.000)
27	(1,1,0,1,0, \mathcal{E})	6.800 (0.551)	4.541 (0.606)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
28	(1,1,0,1,1, \mathcal{E})	6.793 (0.553)	4.532 (0.600)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
29	(1,1,1,0,0, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
30	(1,1,1,0,1, \mathcal{E})	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)	0.000 (0.000)
31	(1,1,1,1,0, \mathcal{E})	6.812 (0.282)	4.567 (0.451)	0.019 (0.577)	0.000 (0.000)	0.000 (0.000)
32	(1,1,1,1,1, \mathcal{E})	6.816 (0.270)	4.620 (0.461)	0.008 (0.584)	0.000 (0.000)	0.000 (0.000)

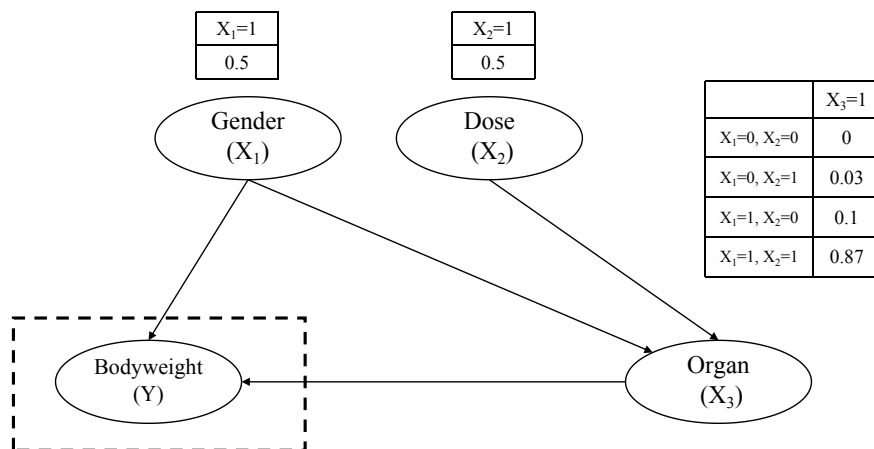


Figure S2: A causal network representing developmental toxicology experiments, including body weight and its potential risk factors.

Table S13: Results of postICEs based on different evidence.

PostICE($Y_{x'} \mid x, Y < 27$)	$(x_1, x_3) = (0, 0)$	$(x_1, x_3) = (0, 1)$	$(x_1, x_3) = (1, 0)$	$(x_1, x_3) = (1, 1)$
$(x'_1, x'_3) = (0, 0)$	0.00	1.59	-7.10	-2.77
$(x'_1, x'_3) = (0, 1)$	-1.42	0.00	-5.50	-3.35
$(x'_1, x'_3) = (1, 0)$	8.81	10.51	0.00	5.58
$(x'_1, x'_3) = (1, 1)$	2.84	4.46	-8.90	0.00

Table S14: Results of posterior causal estimands based on the various evidence for the NTP dataset.

Posterior causal estimands	Evidence $X = x$							
	(0, 0, 0)	(1, 0, 0)	(0, 1, 0)	(1, 1, 0)	(0, 0, 1)	(1, 0, 1)	(0, 1, 1)	(1, 1, 1)
PostNDE($X_1 \Rightarrow Y \mid x, Y < 27$)	8.83	7.00	8.83	7.00	10.00	8.31	10.00	8.12
PostNDE($X_2 \Rightarrow Y \mid x, Y < 27$)	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
PostNDE($X_3 \Rightarrow Y \mid x, Y < 27$)	-1.43	-8.90	-1.43	-8.90	-6.30	-5.51	-6.30	-5.51
PostNIE($X_1 \Rightarrow Y \mid x, Y < 27$)	-0.60	0.00	-5.18	0.00	0.00	-5.51	0.00	-5.30
PostNIE($X_2 \Rightarrow Y \mid x, Y < 27$)	-0.05	-7.58	0.00	0.00	0.00	0.00	-6.30	-4.87
PostNIE($X_3 \Rightarrow Y \mid x, Y < 27$)	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
PostTCE($X_1 \Rightarrow Y \mid x, Y < 27$)	8.23	7.00	3.65	7.00	10.00	2.80	10.00	2.82
PostTCE($X_2 \Rightarrow Y \mid x, Y < 27$)	-0.05	-7.58	0.00	0.00	0.00	0.00	-6.30	-4.87
PostTCE($X_3 \Rightarrow Y \mid x, Y < 27$)	-1.43	-8.90	-1.43	-8.90	-6.30	-5.51	-6.30	-5.51