Causal Additive Models with Unobserved Variables

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Abstract

Causal discovery from data affected by unobserved variables is an important but difficult problem to solve. The effects that unobserved variables have on the relationships between observed variables are more complex in nonlinear cases than in linear cases. In this study, we focus on causal additive models in the presence of unobserved variables. Causal additive models exhibit structural equations that are additive in the variables and error terms. We take into account the presence of not only unobserved common causes but also unobserved intermediate variables. Our theoretical results show that, when the causal relationships are nonlinear and there are unobserved variables, it is not possible to identify all the causal relationships between observed variables through regression and independence tests. However, our theoretical results also show that it is possible to avoid incorrect inferences. We propose a method to identify all the causal relationships that are theoretically possible to identify without being biased by unobserved variables. The empirical results using artificial data and simulated functional magnetic resonance imaging (fMRI) data show that our method effectively infers causal structures in the presence of unobserved variables.

1 INTRODUCTION

A fundamental objective in various fields of science is to identify causal relationships. While randomized control trials are the most effective means of understanding causal relationships, such an approach is often too costly, unethical, or technically impossible to conduct. Thus, causal discovery from purely observational data is very important for scientific research.

Causal discovery methods often assume that the causal structures form directed acyclic graphs (DAGs) and that unobserved common causes are absent [Spirtes and Glymour, 1991, Shimizu et al., 2006, 2011, Hoyer et al., 2009, Mooij et al., 2009, Peters et al., 2014]. If methods that assume the absence of unobserved variables are applied to data affected by unobserved variables, the causal graphs inferred by such methods are biased, and thus tend to be incorrect. The fast causal inference (FCI) [Spirtes et al., 1999] and RFCI [Colombo et al., 2012] both assume the presence of unobserved common causes and can present variable pairs with unobserved common causes. However, they infer causal relationships based on conditional independence, and thus cannot distinguish between causal graphs that entail the same sets of conditional independence.

Until recently, causal functional model-based approaches [Shimizu et al., 2011, Hoyer et al., 2009, Mooij et al., 2009, Zhang and Hyvärinen, 2009, Peters et al., 2011, 2014] had not been used to explore causal models with unobserved variables. Causal functional model-based approaches assume that causal effects can be formulated with a specific form of functions. For example, LiNGAM [Shimizu et al., 2006, 2011] and additive noise models (ANMs) [Hoyer et al., 2009] assume that the data generation process can be formulated as $x_i = f_i(\mathbf{pa}_i) + n_i$, where x_i is an observed variable, pa_i is the set of the direct causes (parents) of x_i , and n_i is the external effect on x_i . These methods identify the causal direction between observed variables x_i and x_j as $x_j \to x_i$ if the residual of x_i regressed on x_j is independent of x_j and the residual of x_i regressed on x_i is dependent of x_i . When analyzing data suited to the models, these approaches can identify the entire causal model.

Recently, a causal functional model-based method called repetitive causal discovery (RCD) [Maeda and Shimizu, 2020], an extension of DirectLiNGAM [Shimizu et al., 2011], was proposed. RCD infers causal graphs in which bi-directed edges represent variable pairs affected by unobserved common causes and directed edges represent the

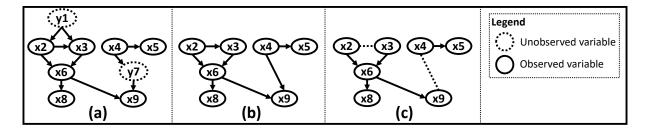


Figure 1: (a) Data generation process. (b) True causal graph of observed variables. (c) Causal graph inferred by our method: Each dashed edge indicates that the causal relationship cannot be determined by our proposed method.

direct causal relationships between observed variables. The RCD method assumes that the causal relationships are linear and the external effects are non-Gaussian. It infers that x_i and x_j have unobserved common causes when the residual of x_i regressed on x_j is dependent of x_j , and vice versa. Janzing et al. [2009] proposed a method to identify the causal relationships between a pair of observed variables assuming that there exists a single unobserved common cause and that the causal functions are nonlinear. However, little research has been conducted on the discovery of causal structures with three or more observed variables, assuming nonlinear causal relationships and the presence of unobserved variables.

The effects that unobserved variables have on the relationships between observed variables are more complex in nonlinear cases than in linear cases. Assume that the causal effect from x_j to x_i is indirectly mediated through unobserved variable y_k (i.e., $x_j \to y_k \to x_i$). Then, $(\exists f, [x_i - f(x_j) \!\perp\!\!\!\perp x_j])$ holds in linear cases but it does not hold in nonlinear cases (i.e., $(\forall f, [x_i - f(x_j) \!\perp\!\!\!\perp x_j]))$. Therefore, the causal relationship between x_i and x_j cannot be determined by regression methods. This is called a cascade ANM (CANM) and has been intensively discussed by Cai et al. [2019]. However, their proposed method assumes that there is no unobserved common cause. Therefore, it is not applicable for inferring causal relationships between three or more observed variables.

Our study is aimed at extending causal additive models (CAMs) [Bühlmann et al., 2014] to incorporate unobserved variables. CAMs are special cases of ANMs, and they assume that the structural equations are additive in the variables and error terms. We call our extended models *causal* additive models with unobserved variables (CAM-UV). In these models, we consider the identifiability of the causal relationships between observed variables. The theoretical results show that it is not possible to identify all the causal relationships, but it is possible to avoid incorrect inferences of causal relationships. We propose a method to infer causal relationships in CAM-UV. Assume that the data generation process is as shown in Figure 1-(a), in which y_1 and y_7 are unobserved variables and the other nodes indicate observed variables. Ideally, the causal graph shown in Figure 1-(b)

should be recovered. However, our goal is to recover the causal graph shown in Figure 1-(c) where the dashed undirected edges between x_2 and x_3 and between x_4 and x_9 indicate that their causal relationships cannot be identified based on our theoretical results.

The contributions of our study are as follows.

- We show the identifiability of the causal relationships between observed variables in *causal additive models* with unobserved variables (CAM-UV).
- We propose a method to infer the causal graph of CAM-UV. Although the method cannot identify all the causal relationships, it can avoid incorrect inferences.
- We provide experimental results on our method and compare them to existing methods using artificial data generated from CAM-UV and simulated functional magnetic resonance imaging (fMRI) data.

All the proofs are available in Maeda and Shimizu [2021].

2 MODEL DEFINITION

Let $X = \{x_i\}$ denote the set of observed variables, $Y = \{y_i\}$ the set of unobserved variables, and $V = \{v_i\}$ the set of all the observed and unobserved variables $(V = X \cup Y)$. We assume the data generation model is formulated as

$$v_i = z_i + w_i + n_i, \quad z_i = \sum_{x_j \in P_i} f_j^{(i)}(x_j), \quad w_i = \sum_{y_k \in Q_i} f_k^{(i)}(y_k)$$

where z_i is the sum of the direct effects of observed variables on v_i , w_i is the sum of the direct effects of unobserved variables on v_i , $f_j^{(i)}$ is a nonlinear function, P_i is the set of observed direct causes of v_i , Q_i is the set of unobserved direct causes of v_i , and n_i is the external effect on v_i . We assume that all the external effects are mutually independent. We also assume that the causal structure of the observed and unobserved variables forms a DAG.

In addition, we impose Assumption 1 (described below) on the causal functions and the external effects in a similar

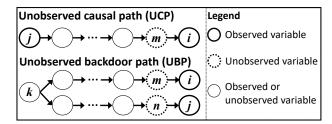


Figure 2: Definitions of an unobserved causal path (UCP) and an unobserved backdoor path (UBP).

way to the Faithfulness assumption [Pearl, 2000, Spirtes et al., 2000]. According to Equation 1, all the observed and unobserved variables are mixtures of external effects generated by the causal functions. In addition, there is an external effect influencing two variables if the two variables have a common *ancestor* (direct or indirect cause) or there is a direct or indirect effect between them. In Assumption 1, we assume that such variables are mutually dependent.

Assumption 1. We assume that all the causal functions and the external effects in CAM-UV satisfy the following condition: If variables v_i and v_j have terms involving functions of the same external effect n_k , then v_i and v_j are mutually dependent (i.e., $(n_k \not\perp v_i) \land (n_k \not\perp v_j) \Rightarrow (v_i \not\perp v_j)$).

3 IDENTIFIABILITY

In this section, we consider the identifiability of the causal relationships between observed variables in CAM-UV.

First, we provide Definitions 1 and 2, which are used in the analysis of the identifiability. The explanatory chart for the definitions is shown in Figure 2.

Definition 1. A directed path from an observed variable to another is called a *causal path* (CP). A CP from x_j to x_i is called an *unobserved causal path* (UCP) if it ends with the directed edge connecting x_i and its unobserved direct cause (i.e., $x_j \to \cdots \to y_m \to x_i$ where y_m is an unobserved direct cause of x_i).

Definition 2. An undirected path between x_i and x_j is called a backdoor path (BP) if it consists of the two directed paths from a common ancestor of x_i and x_j to x_i and x_j (i.e., $x_i \leftarrow \cdots \leftarrow v_k \rightarrow \cdots \rightarrow x_j$, where v_k is the common ancestor). A BP between x_i and x_j is called an unobserved backdoor path (UBP) if it starts with the edge connecting x_i and its unobserved direct cause, and ends with the edge connecting x_j and its unobserved direct cause (i.e., $x_i \leftarrow y_m \leftarrow \cdots \leftarrow v_k \rightarrow \cdots \rightarrow y_n \rightarrow x_j$, where v_k is the common ancestor and v_m and v_m are the unobserved direct causes of v_m and v_m and v_m are the unobserved direct causes of v_m and v_m can be the same variable.

We impose Assumption 2 on the regression functions G_i used in the lemmas provided in this section.

Assumption 2. Let M and N denote sets satisfying $M \subseteq X$ and $N \subseteq X$ where X is the set of all the observed variables in CAM-UV defined in Section 2. We assume that functions $G_i(M)$ take the forms of generalized additive models (GAMs) [Hastie and Tibshirani, 1990] such that $G_i(M) = \sum_{x_m \in M} g_{i,m}(x_m)$ where each $g_{i,m}(x_m)$ is a nonlinear function of x_m . In addition, we assume that functions G_i satisfy the following condition: When both $(x_i - G_i(M))$ and $(x_j - G_j(N))$ have terms involving functions of the same external effect n_k , then $(x_i - G_i(M))$ and $(x_j - G_j(N))$ are mutually dependent (i.e., $(n_k \not \perp x_i - G_i(M)) \land (n_k \not \perp x_j - G_j(N)) \Rightarrow ((x_i - G_i(M)) \not \perp (x_j - G_j(N)))$.

We first show the difference between linear and nonlinear cases of how UCPs and UBPs affect the identifiability of causality. If there is a UCP $x_j \to y_k \to x_i$, then $\exists a \in \mathbb{R}, [x_i - ax_j \perp \!\!\! \perp x_j]$ holds in linear cases [Shimizu et al., 2011] but $\forall g, [x_i - g(x_j) \not \perp x_j]$ holds in nonlinear cases. That is, there is no regression function such that the residual of x_i regressed on x_j is independent of x_j . The observed variable x_i is formulated as $x_i = f_k^{(i)}(f_j^{(k)}(x_j) + n_k) + n_i$. When $f_k^{(i)}$ is a nonlinear function, it cannot be represented as the linear sum of functions of x_j and n_k such as $f_k^{(i)}(f_j^{(k)}(x_j) + n_k) = s(x_j) + t(n_k)$. Therefore, $g(x_j)$ cannot cancel out terms containing x_j from x_i because $g(x_j)$ does not contain n_k . Therefore, when there is a UCP between x_i and x_j , the causal relationship between x_i and x_j cannot be identified through regression and independence tests.

When there is a UBP, there is also a difference in the identifiability of causality between linear and nonlinear cases. In linear cases, the causal relationship between x_i and x_j can be identified if there is a set of observed variables $M\subseteq X\setminus\{x_i,x_j\}$ that blocks all the BPs between x_i and x_j . That is, there exists a variable $x_l\in M$ on each BP [Maeda and Shimizu, 2020]. In nonlinear cases, the causal relationship between x_i and x_j cannot be identified when there is a UBP, regardless of whether it is blocked by observed variables. Let v_m denote the common ancestor of x_i and x_j on a UBP. Because the directed paths from v_m to x_i and to x_j end with their unobserved direct causes, the effect of v_m cannot be removed from x_i or x_j by regression. Therefore, the causal relationship between x_i and x_j cannot be identified when there is a UBP.

In the following, we provide lemmas about the identifiability of causal relationships in CAM-UV. Lemma 1 is about the conditions in which the causal relationship between two observed variables cannot be identified. Lemma 2 is about the condition in which the absence of the direct causal relationship between two observed variables can be identified. Finally, Lemma 3 is about the condition in which the

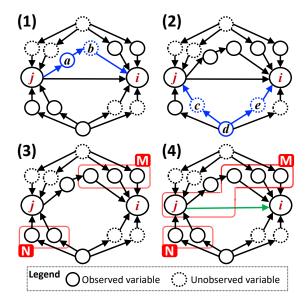


Figure 3: Causal relationships in multivariate cases.

existence and direction of the direct causal relationship between two observed variables can be identified. We provide Examples 1, 2, and 3 for Lemmas 1, 2, and 3, respectively.

Lemma 1. Assume the data generation process of the variables is CAM-UV as defined in Section 2. If and only if Equation 2 is satisfied, there is a UCP or UBP between x_i and x_j where G_1 and G_2 denote regression functions satisfying Assumption 2.

$$\forall G_1, G_2, M \subseteq (X \setminus \{x_i\}), N \subseteq (X \setminus \{x_j\}),$$

$$[(x_i - G_1(M)) \not\perp (x_j - G_2(N))]$$
(2)

Equation 2 indicates that the residual of x_i regressed on any subset of $X \setminus \{x_i\}$ and the residual of x_j regressed on any subset of $X \setminus \{x_j\}$ cannot be mutually independent.

Example 1. In Figure 3-(1), there is a UCP from x_j to x_i (i.e., $x_j \to x_a \to y_b \to x_i$). In Figure 3-(2), there is a UBP between x_j and x_i (i.e., $x_j \leftarrow y_c \leftarrow x_d \to y_e \to x_i$). In these cases, the common effects on x_i and x_j cannot be fully removed by any regression function G_1 or G_2 , because the effects conveyed by the UBP or UCP cannot be removed by G_1 or G_2 .

Lemma 2. Assume the data generation process of the variables is CAM-UV as defined in Section 2. If and only if Equation 3 is satisfied, there is no direct causal relationship between x_i and x_j , and there is no UCP or UBP between x_i and x_j where G_1 and G_2 denote regression functions satisfying Assumption 2.

$$\exists G_1, G_2, M \subseteq (X \setminus \{x_i, x_j\}), N \subseteq (X \setminus \{x_i, x_j\}),$$

$$[((x_i - G_1(M)) \perp \!\!\! \perp (x_j - G_2(N)))]$$

$$(3)$$

Equation 3 indicates that there are regression functions such that the residuals of x_i and x_j regressed on subsets of $X \setminus \{x_i, x_j\}$ are mutually independent.

Example 2. In Figure 3-(3), there is no UCP or UBP between x_i and x_j , and there is no direct causal relationship between x_i and x_j . In Figure 3-(3), M and N are direct causes of x_i and x_j , and they correspond to M and N in Equation 3. They block all the BPs and CPs between x_i and x_j .

Lemma 3. Assume the data generation process of the variables is CAM-UV as defined in Section 2. If and only if Equations 4 and 5 are satisfied, x_j is a direct cause of x_i , and there is no UCP or UBP between x_i and x_j where G_1 and G_2 denote regression functions satisfying Assumption 2.

$$\forall G_1, G_2, M \subseteq (X \setminus \{x_i, x_j\}), N \subseteq (X \setminus \{x_j\}),$$

$$[(x_i - G_1(M)) \not\perp (x_j - G_2(N))]$$
(4)

$$\exists G_1, G_2, M \subseteq (X \setminus \{x_i\}), N \subseteq (X \setminus \{x_i, x_j\}),$$

$$[(x_i - G_1(M)) \perp \!\!\! \perp (x_j - G_2(N))]$$
(5)

Equation 4 indicates that the residual of x_i regressed on any subset of $X \setminus \{x_i, x_j\}$ and the residual of x_j regressed on any subset of $X \setminus \{x_j\}$ cannot be mutually independent. Equation 5 indicates that there are regression functions such that the residual of x_i regressed on a subset of $X \setminus \{x_j\}$ and the residual of x_j regressed on a subset of $X \setminus \{x_i, x_j\}$ are mutually independent.

Example 3. In Figure 3-(4), no UCP or UBP exists between x_j and x_i . There is a direct causal relationship between x_j and x_i . In Figure 3-(4), M and N are direct causes of x_i and x_j , and they correspond to M and N in Equation 5. They block all the BPs and CPs between x_i and x_j including the direct causal effect of x_j on x_i (i.e., $x_j \to x_i$).

Although it is impossible to identify the causal relationship between x_i and x_j when there is a UCP or UBP, it is possible to avoid the incorrect determination of the causal relationship if we use Lemma 1. If there is no UCP or UBP, it is possible to identify the direct causal relationship between x_i and x_j using Lemmas 2 and 3.

Next, we provide Lemma 4, which can be used for identifying a sink of a set of observed variables. Let K denote a set of observed variables. Observed variable x_i is called a sink of K when $x_i \in K$ holds, and each $x_j \in K \setminus \{x_i\}$ is not a descendant of x_i . Example 4 is provided after Lemma 4.

Lemma 4. Assume the data generation process of the variables is CAM-UV as defined in Section 2. Let K denote a set satisfying $K \subseteq X$ and assume $x_i \in K$. If Equation 6 holds, each $x_j \in K \setminus \{x_i\}$ is not a descendant of x_i where G_i^1 , G_j^1 , G_i^2 , and G_j^2 denote regression functions satisfying Assumption 2.

$$\exists G_i^1, M_i \subseteq (X \setminus K),$$

$$\forall x_j \in (K \setminus \{x_i\}), \exists G_j^1, M_j \subseteq (X \setminus K), \forall G_i^2, G_j^2$$

$$[((x_i - G_i^1(M_i \cup K \setminus \{x_i\})) \perp (x_j - G_j^1(M_j)))$$

$$\wedge ((x_i - G_i^2(M_i)) \perp (x_j - G_i^2(M_j)))]$$
(6)

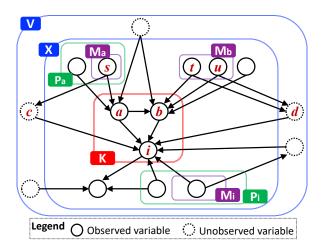


Figure 4: Identification of x_i as a sink of K.

Equation 6 indicates that there exists set $M_i \subseteq (X \setminus K)$ for each $x_i \in K$ satisfying the condition that the residual of x_i regressed on $(M_i \cup K \setminus \{x_i\})$ is independent of the residual of x_i regressed on M_i for each $x_i \in K \setminus \{x_i\}$. In addition, 11 the residual of x_i regressed on M_i cannot be independent of the residual of x_i regressed on M_i for each $x_i \in K \setminus \{x_i\}$.

Example 4. In Figure 4, K consists of three observed variables (i.e., $K = \{x_a, x_b, x_i\}$). The BP between x_i and x_a 13 and the BP between x_i and x_b are blocked by M_a and M_b respectively. In addition, all the effects of x_a and x_b on x_i are mediated by the direct causes of x_i which are included in $K \setminus \{x_i\}$. Then, the residual of x_i regressed on $M_i \cup K \setminus \{x_i\}$ can be independent of the residuals of x_a and x_b regressed on M_a and M_b respectively. In addition, the residual of x_i regressed on M_i cannot be independent of the residuals of x_a and x_b regressed on M_a and M_b respectively. These statements are formulated as Equation 7, which can be generalized to Equation 6.

MODEL ESTIMATION

We propose a method to infer causal relationships between 27 observed variables. The causal graphs inferred by our pro- 28 posed method include directed edges and undirected dashed edges (see Figure 1-(c)). A directed edge indicates a direct 29 causal relationship, and an undirected dashed edge indicates that there is a UCP or UBP between the variables.

First, we propose a method to determine the directed edges. The detailed procedure is listed in Algorithm 1, which consists of two steps. Our method first extracts the candidates

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Algorithm 1: Determine the directed edges
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Input: i.i.d samples of a p-dimensional distribution on
         \{x_1, \cdots, x_p\} X, maximum number of
         variables to examine causality for each step d,
         significance level for independence test \alpha.
Output: the sets of the parents \{M_1, \dots, M_n\}.
function getDirectedEdges(X, d, \alpha)
     # PHASE 1: Extracting the candidates of the
     parents of each variable.
     for i = 1 to p do
         Initialize M_i \leftarrow \emptyset.
     Initialize t \leftarrow 2.
     while t \leq d do
          Initialize noChange \leftarrow True.
         foreach K \in \{K | K \subseteq X, |K| = t\} do
              # Finding the most endogenous variable
              x_b in K
              x_b \leftarrow \operatorname{argmax} \widehat{\mathsf{p-HSIC}}(x_i - G_1(M_i \cup K \setminus
                \{x_i\}), \{x_j - G_2(M_j) | x_j \in K \setminus \{x_i\}\})
              # Computing the independence between
              the residuals in regard to Lemma 4
              e \leftarrow \widehat{\mathsf{p-HSIC}}(x_b - G_1(M_b \cup K \setminus
                \{x_b\}), \{x_j - G_2(M_j) | x_j \in K \setminus \{x_b\}\})
              h \leftarrow \max_{x_j \in K \setminus \{x_b\}} \widehat{\text{p-HSIC}}(x_b -
                G_1(M_b), x_i - G_2(M_i)
              # Checking whether x_b is really a sink of
              if (\alpha < e) \land (\alpha > h) then
                   # When x_b is a sink of K, add each
                   variable in K \setminus \{x_b\} to M_b
                   M_b \leftarrow M_b \cup (K \setminus \{x_b\})
                   noChange \leftarrow False
          # If each M_i remains unchanged, increment t
          by one. If not, substitute 2 for t.
         if noChange = True then
              t \leftarrow t + 1
          else
           t \leftarrow 2
```

PHASE 2: Determining the parents of each variable.

```
for i = 1 to p do
    foreach x_i \in M_i do
         # Checking whether x_i is parent of x_i
         if \alpha < \widehat{\text{p-HSIC}}(x_i - G_1(M_i \setminus \{x_j\}), x_j -
           G_2(M_i)) then
              # When x_i is not a parent, remove it
              from M_i
              M_i \leftarrow M_i \setminus \{x_i\}
return \{M_1, \cdots, M_p\}
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of the parents of each observed variable (lines 2–23 in Algorithm 1), then it determines the parents of each observed variable (lines 24–30). The notations G_1 and G_2 in Algorithms 1 and 2 indicate GAM regression functions. Those functions perform differently in different lines and different iterations.

The first step of Algorithm 1 involves each M_i collecting observed variables that are not descendants of x_i , which we call the candidates of the parents of x_i . The method first initializes each M_i to an empty set (lines 3–4 in Algorithm 1). Then it repeats finding a sink for each K that satisfies $K \subseteq X$ and |K| = t (lines 8–18). That is, each K is a set consisting of t observed variables. The value of t starts at 2 (line 5). It is incremented by 1 when each M_i remains unchanged through an iteration, and it is updated to 2 when at least one M_i changes during the iteration (lines 20–23). When our method determines that x_b is a sink of K, it updates M_b by adding each variable in $K \setminus \{x_b\}$ to M_b (lines 16–17). The iteration ends when t exceeds d (line 6), which is a hyperparameter that is set as the maximum number of |K|. The purpose of d is to reduce the computation time, and it should be set according to the sparsity of the causal relationships.

To find a sink for each K, our proposed method first finds the most endogenous variable x_b in K (lines 9–10). Such a x_b maximizes the independence between $(x_b - G_1(M_b \cup K \setminus \{x_b\}))$ and $(\{x_j - G_2(M_j) | x_j \in K \setminus \{x_b\}\})$. We use the p-value of the Hilbert–Schmidt Independence Criteria (HSIC) [Gretton et al., 2008] for measuring independence, and we also use the GAM regression method proposed by Wood [2004]. Our method examines whether x_b and the other variables in $K \setminus \{x_b\}$ satisfy the condition defined in Lemma 4 using the significance level for independence test, given as hyperparameter α (lines 11–15). If x_b and $K \setminus \{x_b\}$ satisfy the condition, then each variable in $K \setminus \{x_b\}$ is added to M_b (lines 16–17).

In the second step, our proposed method determines the parents of each observed variable. If $x_i \in M_i$ satisfies $x_i - G_1(M_i \setminus \{x_j\}) \perp x_j - G_2(M_j)$, it is not a parent of x_i because of Lemma 2. Therefore, our method removes each x_i satisfying the above equation from M_i and defines the variables remaining in M_i as the parents of x_i (lines 27– 30). The reason why the variables remaining in M_i are the parents of x_i is as follows. Each directed path from each x_i in M_i to x_i is blocked by the parents of x_i that are included in M_i (i.e. $M_i \cap P_i$). If x_i is not a parent of x_i , all the directed paths from x_i to x_i is blocked by $(M_i \cap P_i \setminus \{x_i\})$. Then, $(x_i-G_1(M_i\setminus\{x_i\}))$ and $(x_i-G_2(M_i))$ are mutually independent. If x_i is a parent of x_i , there is a direct causal effect $x_i \to x_i$, and it is not blocked by $(M_i \cap P_i \setminus \{x_i\})$. Then, $(x_i - G_1(M_i \setminus \{x_j\}))$ and $(x_j - G_2(M_j))$ cannot be mutually independent. Therefore, the variables remaining in M_i are parents of x_i .

After determining the direct causal relationships, the pro-

Algorithm 2: Determine the undirected dashed edges

Input: i.i.d samples of a p-dimensional distribution on $\{x_1, \dots, x_p\}$ X, series of the sets of the parents $\{M_1, \dots, M_p\}$, significance level for independence test α .

Output: set of variable pairs having a UCP or UBP C.

1 function getUndirectedEdges $(X, \{M_1, \cdots, M_p\}, \alpha)$ 2 | Initialize $C \leftarrow \emptyset$.

3 | foreach $i, j \ ((1 \le i, j \le p) \land (i \ne j))$ do

4 | if $(x_i \notin M_j) \land (x_j \notin M_i)$ then

5 | if $\alpha > \widehat{\text{p-HSIC}}(x_i - G_1(M_i), x_j - G_2(M_j))$ then

6 | $C \leftarrow C \cup \{\{x_i, x_j\}\}$ 7 | return C

posed method determines variable pairs having UBPs or UCPs (i.e., variable pairs connected with dashed undirected edges). The detailed procedure is listed in Algorithm 2. If the residual of x_i regressed on M_i and that of x_j regressed on M_j are mutually dependent, there is a UCP or UBP between them (lines 5–6 in Algorithm 2). Therefore, our proposed method connects x_i and x_j with a dashed undirected edge.

The time complexity of the method is $\mathcal{O}\left(p2^pn^2\right)$ when d (the maximum number of |K|) equals the number of all the observed variables p (i.e., d=p). Please refer to Maeda and Shimizu [2021] for the details.

5 EXPERIMENTS

We compared the performance of our method to the following methods: PC [Spirtes and Glymour, 1991], FCI [Spirtes et al., 1999], CAM [Bühlmann et al., 2014], RESIT [Peters et al., 2014], and RCD [Maeda and Shimizu, 2020]. PC is a constraint-based method that assumes the absence of unobserved variables. FCI is also a constraint-based method, but it assumes the presence of unobserved variables. CAM and RESIT are causal functional model-based methods that assume that causal functions are nonlinear and unobserved variables are absent. In contrast, RCD is a causal functional model-based method that assumes that causal functions are linear and unobserved variables are present.

The true causal graphs used for the evaluation are defined such that a directed edge is drawn from x_j to x_i when there is a directed path from x_j to x_i on which no other observed variable exists (see Figures 1-(a,b)). There are types of edges other than directed edges (i.e., \leftarrow) in the graphs produced by the above methods and our proposed method, but we only used directed edges for the comparative evaluation.

We used precision, recall, and the F-measure as the evalua-

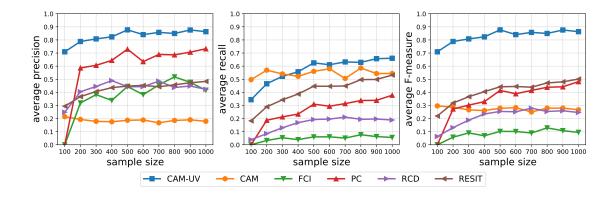


Figure 5: Performance on artificial data generated from causal additive models with unobserved variables.

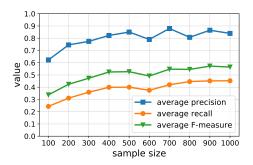


Figure 6: Performance of our method CAM-UV in identifying variable pairs having a UCP or a UBP.

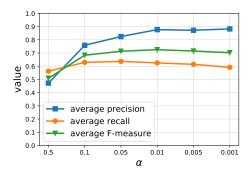


Figure 7: Sensitivity of our method CAM-UV to the setting of α in identifying direct causal relationships.

tion measures. Avoiding false inferences is very important in causal discovery. By evaluating the results in terms of precision, recall, and F-measure, it is possible understand how well each method avoids false inferences. The true positive (TP) is the number of true directed edges that a method correctly infers in terms of their positions and directions. Precision represents the TP divided by the number of estimations, and recall represents the TP divided by the number of all true directed edges. Furthermore, the F-measure is defined as F-measure $= 2 \cdot \operatorname{precision} \cdot \operatorname{recall}/(\operatorname{precision} + \operatorname{recall}).$

We set the significance levels required for the baseline meth-

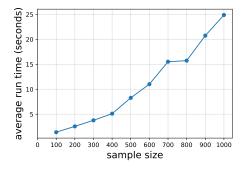


Figure 8: Average runtime of the proposed method.

ods and our proposed method as 0.01. In addition, we set the maximum number of |K| to 3 (i.e., d=3) for our proposed method (see Algorithm 1).

We conducted experiments on the artificial data generated from CAM-UV and the simulated fMRI data created in Smith et al. [2011].

5.1 PERFORMANCE ON ARTIFICIAL DATA GENERATED FROM CAM-UV

Comparison with baseline methods: We performed 100 experiments using artificial data with each sample size $n \in \{100, 200, \cdots, 900, 1000\}$ to compare our method to existing methods. The data for each experiment were generated as follows. The data generation process was accomplished using Equation 1. We prepared ten observed variables, two unobserved common causes, and two unobserved intermediate causal variables. The causal order of the observed variables was determined the same as the order of the indices of the observed variables. The direct causal relationships between the observed variables were determined based on the Erdős–Rényi model [Erdős and Rényi, 1960] with parameter 0.3. That is, each variable pair was connected by an edge with a probability of 0.3. The directions of the edges were determined according to the causal

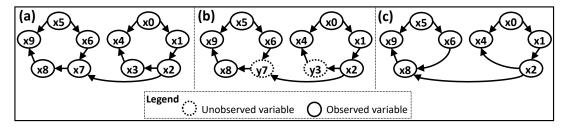


Figure 9: (a) Causal structure in fMRI data (b) Omitted variables. (c) True causal graph after omitting variables.

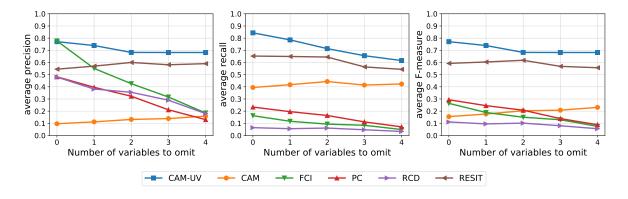


Figure 10: Performance on simulated fMRI data.

order. We drew two directed edges from each unobserved common cause to two randomly selected observed variables. Finally, two variable pairs were randomly chosen, and an unobserved intermediate causal variable was inserted between each variable pair. The indices of the observed variables were randomly permuted after the data were created. The value of each v_i defined in Equation 1 was determined by

$$v_i = \frac{h_i}{\mathrm{sd}(h_i)}, \ h_i = \sum_{v_j \in (P_i \cup Q_i)} ((v_j + a_{i,j})^{c_{i,j}} + b_{i,j}) + n_i,$$
(8)

where $a_{i,j}, b_{i,j}$, and $c_{i,j}$ denote constants, n_i denotes a random variable, and $\mathrm{sd}(h_i)$ denotes the standard deviation of h_i . The values of $a_{i,j}$ and $b_{i,j}$ were randomly chosen from U(-5,5) and U(-1,1), respectively. The value of $c_{i,j}$ was randomly selected from $\{2,3\}$, where the probability of selecting either value is 0.5. The samples of n_i were taken from $U(-10+d_i,10+d_i)$ where d_i is a constant randomly chosen from U(-2,2). The causal effect of v_j on v_i (i.e., $f_j^{(i)}(v_j)$ in Equation 1) corresponds to $((v_j+a_{i,j})^{c_{i,j}}+b_{i,j})/\mathrm{sd}(h_i)$.

Figure 5 shows the results. The graphs plot the mean values of the evaluation measures. CAM-UV scores the best in terms of precision and the F-measure for each sample size. The recall value of CAM-UV increases as the sample size increases. When the sample size is 300 or less, the scores of our proposed method are the second best next to CAM, but it scores the best when the sample size is more than 300.

Performance of identifying UCPs and UBPs: Figure 6 shows how well our proposed method identified UCPs and UBPs. The true positive (TP) is the number of variable pairs having a UCP or UBP and those that are connected by dashed undirected edges in the causal graph inferred by our proposed method. The graphs in Figure 6 show that the precision, recall, and F-measure values increase as the sample size increases.

Sensitivity to the hyperparameter: We conducted experiments to investigate the sensitivity of the proposed method to the settings of the hyperparameter α . We used 500 samples for each experiment with $\alpha \in \{0.5, 0.1, 0.05, 0.01, 0.005, 0.001\}$. Figure 7 shows the results. The precision and F-measure values gradually increase as α decreases, but they remain flat for $\alpha \leq 0.01$.

Average runtime: The average runtimes are shown in Figure 8¹. It was 8.3 seconds when the sample size was 500 and 24.9 seconds when the sample size was 1000.

5.2 PERFORMANCE ON SIMULATED FMRI DATA

We conducted experiments on simulated fMRI data generated by Smith et al. [2011] based on a well-known mathemat-

¹The details of the machine used for computing are as follows. OS: macOS Catalina 10.15.7; CPU: 2.4 GHz 8-core 9th-generation Intel Core i9 processor; Memory: 64 GB 2666 MHz DDR4 memory, Python version: 3.8.6.

ical model of interactions among brain regions, the dynamic causal model [Friston et al., 2003]. We used one of their datasets ("sim2") with ten variables, the causal relationships of which are shown in Figure 9-(a). We randomly omitted m variables for each experiment to create a dataset with unobserved variables. For example, when m=2 and x_3 and x_7 are omitted to make unobserved variables y_3 and y_7 , as shown in Figure 9-(b), the causal graph for evaluation includes directed edges $x_2 \to x_4, x_2 \to x_8$, and $x_6 \to x_8$, as shown in Figure 9-(c). We conducted 100 experiments with 1000 samples randomly extracted from the data for each $m \in \{0,1,2,3,4\}$. Figure 10 shows the results. Though the precision score for CAM-UV is slightly lower than FCI when m=0, our method scores the best for the other cases.

6 CONCLUSIONS

In this study, we extended causal additive models to incorporate unobserved variables, the model for which we called causal additive models with unobserved variables (CAM-UV). Our theoretical analysis showed that the direct causal relationships between observed variables cannot be determined when there is an unobserved causal path (UCP) or unobserved backdoor path (UBP) between the variables. However, the theoretical results also show that it is possible to identify such variable pairs and to avoid incorrect inferences. Based on these theoretical results, we proposed a method to infer causal graphs for CAM-UV and verified the method through experiments. As demonstrated by our theoretical and experimental results, our proposed method is effective in inferring causal relationships in the presence of unobserved variables. Our future research will focus on the application of our method for the efficient intervention design using the results of UBPs and UCPs.

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