# ASPECTS OF GRAPHICAL MODELS CONNECTED WITH CAUSALITY

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#### Abstract

This paper demonstrates the use of graphs as a mathematical tool for expressing independenices, and as a formal language for communicating and processing causal information in statistical analysis. We show how complex information about external interventions can be organized and represented graphically and, conversely, how the graphical representation can be used to facilitate quantitative predictions of the effects of interventions.

We first review the Markovian account of causation and show that directed acyclic graphs (DAGs) offer an economical scheme for representing conditional independence assumptions and for deducing and displaying all the logical consequences of such assumptions. We then introduce the manipulative account of causation and show that any DAG defines a simple transformation which tells us how the probability distribution will change as a result of external interventions in the system. Using this transformation it is possible to quantify, from non-experimental data, the effects of external interventions and to specify conditions under which randomized experiments are not necessary.

Finally, the paper offers a graphical interpretation for Rubin's model of causal effects, and demonstrates its equivalence to the manipulative account of causation. We exemplify the tradeoffs between the two approaches by deriving nonparametric bounds on treatment effects under conditions of imperfect compliance.

### 1 Introduction

Although graphical models are intuitively compelling for conceptualizing statistical associations, the scientific community generally views such models with hesitancy and suspicion. The purpose of this paper is to demonstrate the use of graphs as a precise mathematical tool of great versatility, especially as a formal language for communicating causal information in statistical analysis.

Causal models, no matter how they are represented, discovered, or tested, are generally regarded as more useful than associational models because causal models provide information about the dynamics of the system under study. In other words, a joint distribution tells us how probable events are and how probabilities would change with subsequent observations, but a causal model also tells us how these probabilities would change as a result of external interventions in the system. For this reason, causal models (or "structural models" as they are often called) have been the target of relentless scientific pursuit and, at the same time, the center of much controversy and speculation [Freedman 1987].

The directionality of arcs in graphical models has been treated very cautiously in the statistical literature [Lauritzen & Spiegelhalter 1988, Cox 1992, Cox & Wermuth (in press), Spiegelhalter etal. (in press)]: the causal interpretation of the directed arcs has been de-emphasized in favor of the safer interpretation in terms of "relevance" and "dependence". This limited interpretation is deficient in several respects. First, causal associations are the primary source of judgments about dependence and relevance; they should therefore guide preformal thinking about the design of statistical studies [Dempster 1990]. Second, rejecting the causal interpretation of directed arcs prevents us from using graphical models for making legitimate predictions about the effects of actions. Such predictions are indispensable in most decision making applications, including policy analysis and treatment management.

The primary aim of this paper is to show how complex information about external interventions can be organized and represented graphically and, conversely, how the graphical representation can be used to facilitate quantitative predictions of the effects of interventions.

Section 2 will review the use of directed acyclic graphs (DAGs) as a language for communicating conditional independence assumptions. Sections 3 and 4 will define the causal interpretation of DAGs and Section 5 will demonstrate their use in observational studies. Section 6 will demonstrate the equivalence between the language of graphs and Rubin's model of causal effects. Finally, Section 7 applies the two approaches to the analysis of treatment effects in experimental studies with imperfect compliance. Using this example we show how a latent-variable structure can be reduced to an equivalent counterfactual model and how the two approaches can be used to derive nonparametric bounds on the causal effects of treatments, when data is taken under conditions of partial compliance.

# 2 Directed Graphs and Conditional Independence: A Review

Networks employing directed acyclic graphs (DAGs) are primarily used to provide either

- 1. an economical scheme for representing conditional independence assumptions, or
- 2. a graphical language for representing causal influences.

This section will focus on the former, since causal influences are discussed in the remaining parts of this paper.

Given a DAG  $\Gamma$  and a joint distribution P over a set  $X = \{X_1, ..., X_n\}$  of discrete variables, we say that  $\Gamma$  represents P if there is a one-to-one correspondence between the variables in X and the nodes of  $\Gamma$ , such that P admits the recursive product decomposition

$$P(x_1, ..., x_n) = \prod_i P(x_i \mid \mathbf{pa}_i)$$
 (1)

where  $\mathbf{pa}_i$  are the direct predecessors (called *parents*) of  $X_i$  in  $\Gamma$ . For example, the DAG in Figure 1 induces the decomposition

$$P(x_1, x_2, x_3, x_4, x_5) = P(x_1) P(x_2|x_1) P(x_3|x_1) P(x_4|x_2, x_3) P(x_5|x_4)$$
 (2)

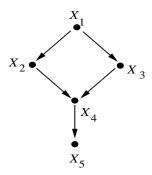


Figure 1: A typical directed acyclic graph (DAG) representing the decomposition of Eq. (2).

The recursive decomposition in Eq. (1) implies that, given its parent set  $\mathbf{pa}_i$ , each variable  $X_i$  is conditionally independent of all its other predecessors  $\{X_1, X_2, ..., X_{i-1}\}\setminus \mathbf{pa}_i$ . Using Dawid's [1980] notation, we can state this set of independencies as follows:

$$X_i \parallel \{X_1, X_2, ..., X_{i-1}\} \backslash \mathbf{pa}_i \mid \mathbf{pa}_i \quad i = 2, ..., n$$
 (3)

Such a set of independencies will be called *Markovian*, since it reflects the Markovian condition for state transitions: Each state is rendered independent of the past, given

its immediately preceding state. For example, the DAG of Figure 1 implies the following Markovian independencies:

$$X_2 \parallel \{0\} \mid X_1, \quad X_3 \parallel X_2 \mid X_1, \quad X_4 \parallel X_1 \mid \{X_2, X_3\}, \quad X_5 \parallel \{X_1, X_2, X_3\} \mid X_4 \parallel X_5 \parallel$$

Conversely, any list M of Markovian independencies identifies a DAG  $\Gamma$  (that represents P) because it permits a recursive product decomposition as in Eq. (1). However, such decomposition may imply additional independencies that are not included in M. For example, the decomposition of Eq. (1) implies  $X_5 \parallel X_3 \mid \{X_1, X_4\}$  (which is not part of M) regardless of the numerical values assigned to the factors of that product. A graphical criterion called d-separation [Pearl 1988] permits us to read off the DAG the sum total of all independencies implied by a given decomposition.

**Definition 2.1** (*d-separation*) If X, Y, and Z are three disjoint subsets of nodes in a DAG  $\Gamma$ , then Z is said to *d*-separate X from Y, denoted  $d(X, Z, Y)_{\Gamma}$ , if and only if there is no path from a node in X to a node in Y along which the following two conditions hold: (1) every node with converging arrows either is or has a descendant in Z, and (2) every other node is outside Z. A path satisfying the conditions above is said to be *active*; otherwise it is said to be *blocked* (by Z). By *path* we mean a sequence of consecutive edges (of any directionality) in the DAG.

In Figure 1, for example,  $X = \{X_2\}$  and  $Y = \{X_3\}$  are d-separated by  $Z = \{X_1\}$ ; the path  $X_2 \leftarrow X_1 \rightarrow X_3$  is blocked by  $X_1 \in Z$ , while the path  $X_2 \rightarrow X_4 \leftarrow X_3$  is blocked because  $X_4$  and all its descendants are outside Z. Thus  $d(X_2, X_1, X_3)$  holds in  $\Gamma$ . However, X and Y are not d-separated by  $Z' = \{X_1, X_5\}$  because the path  $X_2 \rightarrow X_4 \leftarrow X_3$  is rendered active by virtue of  $X_5$ , a descendant of  $X_4$ , being in Z. Consequently,  $d(X_2, \{X_1, X_5\}, X_3)$  does not hold in  $\Gamma$ ; Metaphorically, learning the value of the consequence  $X_5$  renders its causes  $X_2$  and  $X_3$  dependent, as if a pathway were opened along the converging arrows at  $X_4$ .

The d-separation criterion has been shown to be both sound and complete relative to the set of distributions that are represented by a DAG  $\Gamma$  [Verma 1986, Geiger & Pearl 1988]. In other words, there is a one-to-one correspondence between the set of independencies implied by the recursive decomposition of Eq. (1) and the set of triples (X, Z, Y) that satisfy the d-separation criterion in  $\Gamma$ . Furthermore, the d-separation criterion can be tested in time linear in the number of edges in  $\Gamma$ . Thus, a DAG can be viewed as an efficient scheme for representing Markovian independence assumptions and for deducing and displaying all the logical consequences of such assumptions. Additional properties of DAGs and their applications to evidential reasoning in expert systems are discussed in [Pearl 1988, Pearl et al. 1990, Geiger 1990, Lauritzen & Spiegelhalter 1988, Spiegelhalter et al. (in press), Pearl 1993a].

# 3 Graphical Models and the Manipulative Account of Causation

The interpretation of DAGs as carriers of independence assumptions does not specifically mention causation and will in fact be valid for any set of Markovian independencies, along any ordering (not necessarily causal or chronological) of the variables.

However, the patterns of independencies portrayed in a DAG are so typical of causal organizations that some of these patterns can only be given meaningful interpretation in terms of causation. For example, we can hardly find a pair of dependent events,  $E_1$  and  $E_2$ , that are rendered independent by conditioning on a third event  $E_3$  unless  $E_3$  serves as a cause for either  $E_1$  or  $E_2$  (or both). Indeed, we cannot even conceive of three such events if we constrain  $E_3$  to occur after  $E_1$  and  $E_2$ , so as to suppress the causal interpretation above. The DAG representation provides a perfect language for such dependencies; it lets  $E_3$  d-separate  $E_2$  from  $E_1$  in the pattern  $E_1 \to E_3 \to E_2$  or  $E_1 \leftarrow E_3 \to E_2$ , but not in the converging pattern  $E_1 \to E_3 \leftarrow E_2$ . This distinction is the basis for the Markovian accounts of causation, as exemplified by those of [Granger 1988, Suppes 1970], and by the more elaborate, non-temporal accounts of [Pearl & Verma 1991] and [Spirtes etal. 1993].

However, the Markovian account still leaves open the question of why such intricate patterns of independencies are produced by and become the characteristic signature of causal organizations. A related question is how these patterns are connected with the more basic notions associated with causation, such as influence, manipulation, and control. The connection is made in the mechanism-based account of causation.

The basic idea behind this account goes back to [Simon 1977] and is stated succinctly in his forward to [Glymour et al. 1987]: "The advantage of representing the system by structural equations that describe the direct causal mechanisms is that if we obtain some knowledge that one or more of these mechanisms has been altered, we can use the remaining equations to predict the consequences – the new equilibrium." Here, by "mechanism" Simon means any stable relationship between two or more variables, usually expressed in functional form, that remains invariant to external influences until it falls directly under such influences.

This mechanism-based model was adapted in [Pearl & Verma 1991] for defining probabilistic causal theories; each child-parent family in a DAG  $\Gamma$  represents a deterministic function  $X_i = f_i(\mathbf{pa}_i, \epsilon_i)$ , where  $\mathbf{pa}_i$  are the parents of variable  $X_i$  in  $\Gamma$ , and  $\epsilon_i$ , 0 < i < n, are mutually independent, arbitrarily distributed random disturbances such that each  $\epsilon_i$  is independent on all non-descendants of  $X_i$ . Characterizing each child-parent relationship as a deterministic function, instead of the usual conditional probability  $P(x_i \mid \mathbf{pa}_i)$ , imposes equivalent independence constraints on the resulting distributions and leads to the same recursive decomposition

$$P(x_1, ..., x_n) = \prod_i P(x_i \mid \mathbf{pa}_i)$$
 (5)

that characterizes DAG models (see Eq. 1). However, the functional characterization  $X_i = f_i(\mathbf{pa}_i, \epsilon_i)$  also specifies how the resulting distribution would change in response to external interventions, since, by convention, each function is presumed to remain constant unless specifically altered. Moreover, the non-linear character of  $f_i$  permits us to treat changes in the function  $f_i$  itself as a variable,  $F_i$ , by writing

$$X_i = f_i'(\mathbf{pa}_i, F_i, \epsilon_i) \tag{6}$$

where

$$f'_i(a, b, c) = f_i(a, c)$$
 whenever  $b = f_i$ .

Thus, any external intervention  $F_i$  that alters  $f_i$  can be represented graphically as an added parent node of  $X_i$ , and the effect of such an intervention can be analyzed by Bayesian conditionalization, that is, by simply setting this added parent variable to the appropriate value  $f_i$ .

The simplest type of external intervention is one in which a single variable, say  $X_i$ , is forced to take on some fixed value  $x'_i$ . Such intervention, which we call atomic, amounts to replacing the old functional mechanism  $X_i = f_i(\mathbf{pa}_i, \epsilon_i)$  with a new mechanism  $X_i = x'_i$  governed by some external force  $F_i$  that sets the value  $x'_i$ . If we imagine that each variable  $X_i$  could potentially be subject to the influence of such an external force  $F_i$ , then we can view the causal network  $\Gamma$  as an efficient code for predicting the effects of atomic interventions and of various combinations of such interventions.

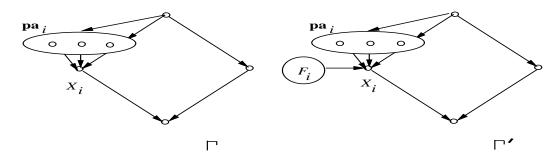


Figure 2: Representing external intervention  $F_i$  by an augmented network  $\Gamma' = \Gamma \cup \{F_i \to X_i\}.$ 

The effect of an atomic intervention  $set(X_i = x_i')$  is encoded by adding to  $\Gamma$  a link  $F_i \longrightarrow X_i$  (see Figure 2), where  $F_i$  is a new variable taking values in  $\{set(x_i'), idle\}$ ,  $x_i'$  ranges over the domain of  $X_i$ , and idle represents no intervention. Thus, the new parent set of  $X_i$  in the augmented network is  $\mathbf{pa}_i' = \mathbf{pa}_i \cup \{F_i\}$ , and it is related to  $X_i$  by the conditional probability

$$P(x_i \mid \mathbf{pa}_i') = \begin{cases} P(x_i \mid \mathbf{pa}_i) & \text{if } F_i = idle \\ 0 & \text{if } F_i = set(x_i') \text{ and } x_i \neq x_i' \\ 1 & \text{if } F_i = set(x_i') \text{ and } x_i = x_i' \end{cases}$$

$$(7)$$

The effect of the intervention  $set(x_i')$  is to transform the original probability function  $P(x_1,...,x_n)$  into a new function  $P_{x_i'}(x_1,...,x_n)$ , given by

$$P_{x_i'}(x_1, ..., x_n) = P'(x_1, ..., x_n \mid F_i = set(x_i'))$$
(8)

where P' is the distribution specified by the augmented network  $\Gamma' = \Gamma \cup \{F_i \to X_i\}$  and Eq. (7), with an arbitrary prior distribution on  $F_i$ . In general, by adding a hypothetical intervention link  $F_i \to X_i$  to each node in  $\Gamma$ , we can construct an augmented probability function  $P'(x_1, ..., x_n; F_1, ..., F_n)$  that contains information about richer types of interventions. Multiple interventions would be represented by conditioning P' on a subset of the  $F_i$ 's (taking values in their respective  $set(x_i')$ ), while the pre-intervention probability function P would be viewed as the posterior distribution induced by conditioning each  $F_i$  in P' on the value idle.

Eq. (8) explains why randomized experiments are sufficient for estimating the effect of interventions even when the causal network is not given: because the intervening variable  $F_i$  enters the networks as a root node (i.e., independent of all other ancestors of  $X_i$ ) it is equivalent to a treatment-selection policy governed by a random device.

### 4 A Transformation Formula for Interventions

This representation yields a simple and direct transformation between the pre-intervention and the post-intervention distributions:

$$P_{x_i'}(x_1, ..., x_n) = \begin{cases} \frac{P(x_1, ..., x_n)}{P(x_i \mid \mathbf{pa}_i)} & \text{if } x_i = x_i' \\ 0 & \text{if } x_i \neq x_i' \end{cases}$$
(9)

This transformation reflects the removal of the term  $P(x_i | \mathbf{pa}_i)$  from the product decomposition of Eq. (5), since  $\mathbf{pa}_i$  no longer influence  $X_i$ . Graphically, the removal of this term is equivalent to removing the links between  $\mathbf{pa}_i$  and  $X_i$ , while keeping the rest of the network intact. Transformations involving conjunctive and disjunctive actions can be obtained by straightforward applications of Eq. (8) [Spirtes et al. 1993, Goldszmidt & Pearl 1992, Goldszmidt 1992]

The transformation (9) exhibits the following properties:

- 1. An intervention  $set(x_i)$  can affect only the descendants of  $X_i$  in  $\Gamma$ .
- 2. For any set **S** of variables, we have

$$P_{x_i'}(\mathbf{S} \mid \mathbf{pa}_i) = P(\mathbf{S} \mid x_i', \mathbf{pa}_i). \tag{10}$$

In other words, given  $X_i = x_i'$  and  $\mathbf{pa}_i$ , it is superfluous to find out whether  $X_i = x_i'$  was established by external intervention or not. This can be seen directly from the augmented network  $\Gamma'$  (see Figure 2), since  $\{X_i\} \cup \mathbf{pa}_i$  d-separates  $F_i$  from the rest of the network, thus legitimizing the conditional independence  $\mathbf{S} \parallel F_i \mid (X_i, \mathbf{pa}_i)$ .

3. A necessary and sufficient condition for an external intervention  $set(X_i = x_i')$  to have the same effect on  $X_j$  as the passive observation  $X_i = x_i'$  is that  $X_i$  d-separates  $\mathbf{pa}_i$  from  $X_j$ , that is,

$$P_{x_i'}(x_j) = P(x_j \mid x_i') \quad \text{iff} \quad X_j \parallel \mathbf{pa}_i \mid X_i. \tag{11}$$

The immediate implication of Eq. (9) is that, given the structure of the causal network  $\Gamma$ , one can infer post-intervention distributions from pre-intervention distributions; hence, we can reliably estimate the effects of interventions from passive (i.e., non-experimental) observations. Of course, Eq. (9) does not imply that we can always substitute observational studies for experimental studies, as this would require an estimation of  $P(x_i \mid \mathbf{pa}_i)$ . The mere identification of  $\mathbf{pa}_i$  (i.e., the direct causal

factors of  $X_i$ ) requires substantive causal knowledge of the domain which is often unavailable. Moreover, even when we have sufficient substantive knowledge to structure  $\Gamma$ , some members of  $\mathbf{pa}_i$  may be unobservable, or *latent*. Fortunately, there are conditions for which an unbiased estimate of  $P_{x_i'}(x_j)$  can be obtained even when the  $\mathbf{pa}_i$  variables are latent and, moreover, a simple graphical criterion can tell us when these conditions are satisfied.

# 5 Controlling Confounding Bias: The Back-door Criterion

Assume we are given a causal network  $\Gamma$  together with non-experimental data on a subset  $\mathbf{X}_o$  of observed variables in  $\Gamma$  and we wish to estimate what effect the intervention  $set(X_i = x_i')$  would have on some response variable  $X_j$ . In other words, we seek to estimate  $P_{x_i'}(x_j)$  from a sample estimate of  $P(\mathbf{X}_o)$ . Applying Eq. (8), we can write

$$P_{x'_{i}}(x_{j}) = P'(x_{j} \mid F_{i} = set(x'_{i}))$$

$$= \sum_{\mathbf{S}} P'(x_{j} \mid \mathbf{S}, X_{i} = x'_{i}, F_{i} = set(x'_{i}))P'(\mathbf{S} \mid F_{i} = set(x'_{i}))$$
(12)

where **S** is any set of variables. Clearly, if **S** satisfies

$$\mathbf{S} \parallel F_i \text{ and } X_j \parallel F_i \mid (X_i, \mathbf{S})$$
 (13)

then Eq. (12) can be reduced to

$$P_{x'_{i}}(x_{j}) = \sum_{\mathbf{S}} P(x_{j} \mid \mathbf{S}, x'_{i}) P(\mathbf{S})$$

$$= E_{\mathbf{S}}[P(x_{j} \mid \mathbf{S}, x'_{i})]$$
(14)

Thus, if we find a set  $\mathbf{S} \subseteq \mathbf{X}_o$  of observables satisfying Eq. (13), we can estimate  $P_{x_i'}(x_j)$  by taking the expectation (over  $\mathbf{S}$ ) of  $P(x_j \mid \mathbf{S}, x_i')$ , and the latter can easily be estimated from non-experimental data. It is also easy to verify that Eq. (13) is satisfied by any set  $\mathbf{S}$  that meets the following  $back-door^1$  criterion:

- 1. No node in **S** is a descendant of  $X_i$ , and
- 2. **S** d-separates  $X_i$  from  $X_j$  along every path containing an arrow into  $X_i$ .

In Figure 3, for example, the sets  $\mathbf{S}_1 = \{X_3, X_4\}$  and  $\mathbf{S}_2 = \{X_4, X_5\}$  meet the back-door criterion, but  $\mathbf{S}_3 = \{X_4\}$  does not because  $X_4$  does not d-separate  $X_i$  from  $X_j$  along the path  $(X_i, X_3, X_1, X_4, X_2, X_5, X_j)$ . Thus, we have obtained a simple graphical criterion for finding a set of observables for estimating (by conditioning) the effect of interventions from purely non-experimental data.

<sup>&</sup>lt;sup>1</sup>The name "back-door" echoes condition 2, which requires that only indirect paths from  $X_i$  to  $X_j$  be d-separated; these paths can be viewed as entering  $X_i$  through the back door.

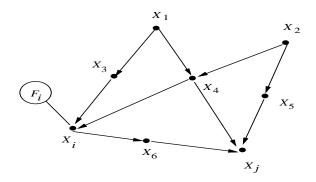


Figure 3: A DAG representing the back-door criterion; adjusting for variables  $\{X_3, X_4\}$  (or  $\{X_4, X_5\}$ ) yields an unbiased estimate of  $P(x_j \mid set(x'_i))$ .

It is interesting that the conditions formulated in Eq. (13) are equivalent to those known as strongly ignorable treatment assignment (SITA) conditions in Rubin's model for causal effect [Rosenbaum & Rubin 1983] (see Section 6 for detailed comparison). Reducing the SITA conditions to the graphical back-door criterion facilitates the search for an optimal conditioning set **S** and significantly simplifies the judgments required for ratifying the validity of such conditions in practical situations.

Eq. (9) was derived under the assumption that the pre-intervention probability P is given by the product of Eq. (5), which represents a joint distribution prior to making any observations. To predict the effect of action  $F_i$  after observing C, we must also invoke assumptions about persistence, so as to distinguish properties that will terminate as a result of  $F_i$  from those that will persist despite  $F_i$ . Such a model of persistence was invoked in [Pearl 1993b]; there, it was assumed that only those properties that are not under any causal influence to terminate should persist. This assumption yields formulas for the effect of conditional interventions (conditioned on the observation C). Again, given  $\Gamma$ , these effects can be estimated from non-experimental data.

[Spirtes et al. 1993] have explored a more ambitious task – estimation of the effect of intervention when the structure of  $\Gamma$  is not available and must also be inferred from the data. Recent developments in graphical models [Pearl & Verma 1991, Spirtes et al. 1993] have produced methods that, under certain conditions, permit us to infer plausible causal structures from non-experimental data, albeit such structures have a weaker set of guarantees than those obtained through controlled randomized experiments. These guarantees fall into two categories: minimality and stability [Pearl & Verma 1991]. Minimality guarantees that any other causal structure compatible with the data is necessarily more redundant, and hence less trustworthy, than the one(s) inferred. Stability ensures that any alternative structure compatible with the data must be less stable than the one(s) inferred; namely, slight fluctuations in the distributions of the disturbances  $\epsilon_i$  (Eq. (6)) will render an alternative structure no longer compatible with the data.

When the structure of  $\Gamma$  is to be inferred under these guarantees, the formulas governing the effects of interventions and the conditions required for estimating these effects become rather complex [Spirtes et al. 1993]. Alternatively, one can produce

bounds on the effects of interventions by taking representative samples of inferred structures and estimating  $P_{x'_i}(x_j)$  according to Eq. (9) (or Eq. (14)) for each such sample.

### 6 Relation to Rubin's Model of Causal Effects

So far, our discussion of causal graphs has focused on the manipulative account of causation which, as was shown in Section 3, coincides with Simon's mechanism-based account. Another view of these basic accounts is provided by the counterfactual model developed by [Rubin 1974] and [Holland 1986, Rosenbaum & Rubin 1983, Pratt & Schlaifer 1988], the roots of which date back to [Neyman 1935] and [Fisher 1935].

In Rubin's model, we imagine that an intervention Z (or "treatment" as it is often called) can be applied at various levels 1, 2, ..., T to any experimental subject (called a "unit") and that it is possible to record the values of the response observed in conjunction with the different levels of the treatment. The correspondence between the applied levels of the treatment and the recorded levels of the response would then constitute the "causal effect" associated with the particular subject, as it characterizes the potential impact of the treatment if applied (counterfactually) to that subject. The target of causal-inference analysis is then the estimation, from statistical data, of the properties of the causal effect vector  $\mathbf{r} = (r_1, r_2, ..., r_t, ..., r_T)$ , where  $r_t$  stands for the response that the subject would exhibit if the t-th level of the treatment were applied. For any given subject,  $r_t$  is considered a deterministic (albeit unobservable) entity, as it determines precisely the response of the subject, had he/she been given the treatment Z = t. However, for a subject randomly drawn from a population, we can view  $r_t$  as a random variable, and therefore, we can attempt to estimate its distribution, its expectation, or the expectation of the difference  $r_t - r_{t'}$ .

The distribution of  $r_t$ , using the language of the manipulative account, is equal to the distribution of the observed response Y, conditioned on the intervention  $F_Z = set(Z = t)$ , namely,

$$P(r_t = y) = P'(Y = y|set(Z = t)) = P_{Z=t}(y)$$
(15)

and

$$E(r_t - r_{t'}) = E(Y|set(Z = t)) - E(Y|set(Z = t'))$$

$$\tag{16}$$

The reason we must condition on the action  $F_Z = set(Z = t)$  and not on the observation Z = t is that, to comply with the interpretation of  $r_t$  as the subject's hypothetical response to treatment Z = t, we must suppress any information that the assignment Z = t may provide on the nature of subject.

The translation provided by Eqs. (15)-(16) implies that the causal effect defined as  $E(r_t-r_{t'})$  can be computed from the manipulative account defined in Section 3 and its associated transformations, as given in Eq. (9). This translation also permits us to devise a graphical representation to Rubin's model, thus displaying the functional role of  $r_t$ . For example, if in Figure 3 we take  $X_i$  to be the treatment variable Z and  $X_j$  to be the observed response Y, then the graph associated with Rubin's

model would correspond to the one in Figure 4. The arc from  $X_3$  to Z represents a non-randomized treatment assignment policy, where the assignment of subjects to treatment Z may depends on the factor  $X_3$ . The main difference between the two

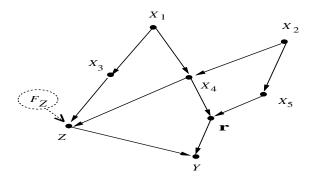


Figure 4: Graphical representation of Rubin's model, showing the observed response (Y) as a function of treatment (Z) and causal effect  $(\mathbf{r})$  variables.

figures is that in the counterfactual model of Figure 4  $\mathbf{r}$  is treated explicitly as a variable, whereas in the manipulative model of Figure 3  $\mathbf{r}$  is represented implicitly as a function that connects Y to its direct causal factors:  $X_4, X_5$ , and  $X_6$ . The two alternative representations of  $\mathbf{r}$  are in line with the transformation defined in Eq. (6). Note also that the counterfactual reading of  $\mathbf{r}$  is an integral part of the mechanism-based reading of causation; the semantics of the function  $f_i(\mathbf{pa}_i, \epsilon_i)$  is intrinsically counterfactual because it defines the value of  $X_i$  for any hypothetical value combination of  $\mathbf{pa}_i$  and  $\epsilon_i$ .

It is not hard to verify that Figures 3 and 4 are empirically equivalent, in the sense that they imply the same statistical and manipulative behavior for all observed variables. For example, our back-door criterion between  $X_i$  and  $X_j$  (see Eq. (13)) translates to an equivalent back-door criterion between Z and  $\mathbf{r}$ ,

$$Z \parallel \mathbf{r} \mid \mathbf{S}$$
 (17)

This is precisely the SITA condition defined in [Rosenbaum & Rubin 1983]. Moreover, since Dawid's [1980] axioms for conditional independence are faithfully encoded in the d-separation criterion, we can immediately translate the condition in Eq. (17) into equivalent graphical criteria, all of which are vividly displayed in the graph. For example, stated in terms of the unobserved set of variables  $\mathbf{U}$  ( $\mathbf{U} = \{X_1, X_2\}$  in Figure 4), our back-door criterion (Eq. (17)) reads

$$Z \parallel \mathbf{U} \mid \mathbf{S} \text{ or } \mathbf{U} \parallel \mathbf{r} \mid \mathbf{S}$$
 (18)

These are precisely the alternative conditions for (X, U)-adjustable treatment assignment given in [Rosenbaum 1989].

The main attraction of Rubin's model is that it allows us to precisely define the causal quantities we wish to estimate without specifying the inference methods used in obtaining these estimates. As a result, the model exposes the fundamental assumptions needed to make the desired estimates feasible, and we are often able to reduce these assumptions to statements about independencies which, at least in principle, can be submitted to judgmental verification.

Since quantities defined in Rubin's model can be translated to equivalent quantities in the manipulative account of causation (see Eqs. (15)-(16)), it is clear that the latter should enjoy similar advantages. Moreover, considering that graphical models provide a calculus for processing manipulative statements (through the introduction of hypothetical action variables, as shown in Section 3), it is not surprising that graphical techniques are applicable for processing statements articulated in Rubin's model.

The current popularity of Rubin's model is in part a reaction to basic inadequacies of the structural equations framework, which forces the analyst to commit to a particular regression model, governed by a particular set of random variables, *iid* disturbances, and hypothetical parameters. While the graphical framework indeed commits the analyst to treating quantities as random variables, often latent, it does not require any assumption of *iid* or parametric structure. The analyst is committed *only* to the qualitative structure behind causal thinking which, we conjecture, is the very structure an analyst must consult when judging the assumptions stated in Rubin's model.

It is not surprising, then, that the two approaches yield identical conclusions in all cases where such conclusions can be stated formally or tested empirically. In cases where the conclusions involve human judgment (e.g., confirming the SITA conditions), the two approaches provide complementary languages for phrasing the judgments required; only time will tell under what circumstances one language will be deemed more natural than the other.

# 7 Example: Causal Effects Under Partial Compliance

To demonstrate the interplay between the counterfactual and the latent-variable models, we will present an analysis of a well-known practical problem using the two approaches.

### 7.1 The problem

Consider an experimental study in which random assignment has taken place but compliance is not perfect, that is, the treatment received is different from that assigned. It is well known that under such conditions a bias may be introduced, in the sense that the true causal effect of the treatment may deviate substantially from the causal effect computed by simply comparing subjects receiving the treatment with those not receiving the treatment. Thus, for example, subjects who did not comply with the assigned treatment may be precisely those who would have responded adversely to the treatment, so the treatment, when applied uniformly to the population, might actually be substantially less effective than the study reveals.

In an attempt to compensate for such bias, economists have devised correctional formulas, called "instrumental variables" [Bowden & Turkington 1984], which, in general, do not hold outside the linear regression model. A recent analysis by Efron and Feldman [1991] represents a healthy departure from the linear regression model, yet it still makes restrictive commitments to a particular mode of interaction between compliance and response. Angrist et al. [1993], invoking Rubin's model, have identified a set of assumptions under which the "instrumental variable" formula is valid, but have not provided an alternative, assumption-free formula. We now derive correctional formulas that rely solely on observed quantities and are universal, that is, they are valid no matter what model actually governs the interactions between compliance and response.

#### 7.2 The latent-structure approach

The canonical partial-compliance setup can be represented by the following network:

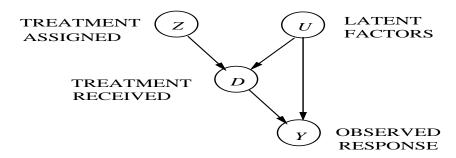


Figure 5: Graphical representation of causal dependencies in a randomized clinical trial with partial compliance.

We assume that Z, D, and Y are observed binary variables where, using conventional terminology (e.g., [Angrist et al. 1993]), Z represents the (randomized) "treatment assignment", D is the treatment actually received, and Y is the observed response. U represents all unobserved and unknown factors which, as Figure 5 shows, may influence the outcome Y and the treatment D. To simplify the notation, we let z, d, and y represent, respectively, the values taken by the variables Z, D, and Y, with the following interpretation:

 $z \in \{z_0, z_1\}, z_1$  asserts that treatment has been assigned  $(z_0, its negation)$ 

 $d \in \{d_0, d_1\}, d_1$  asserts that treatment has been administered  $(d_0, \text{ its negation})$ 

 $y \in \{y_0, y_1\}, y_1$  asserts a positive observed response  $(y_0, \text{ its negation})$ 

The domain of U remains unspecified and may, in general, combine the spaces of several random variables, both discrete and continuous.

The graphical model reflects two assumptions:

1. The treatment assignment does not influence Y directly, only through the actual treatment D, that is,

$$Z \parallel Y \mid \{D, U\} \tag{19}$$

In practice, any direct effect Z might have on Y would be adjusted for through the use of a placebo.

2. Z and U are marginally independent, that is,  $Z \parallel U$ . This independence is partly ensured through the randomization of Z, which rules out any common cause for both Z and U. The absence of a direct path from Z to U represents the assumption that latent factors which affect compliance with the assignment are not in themselves affected by the assignment.

These assumptions impose the following decomposition on the joint distribution

$$P(y, d, z, u) = P(y|d, u) P(d|z, u) P(z) P(u)$$
(20)

which, of course, cannot be observed directly. However, the marginal distribution P(y,d,z) and, in particular, the conditional distribution  $P(y,d|z), z = z_0, z_1$  are observed, and the challenge is to estimate the causal effect of D on Y from these distributions.

For any two binary variables X and Y, define the causal effect  $R(X \to Y)$  of X on Y as

$$R(X \to Y) = P(y_1|set(x_1)) - P(y_1|set(x_0))$$
 (21)

Thus, for the experimental design depicted in Figure 5, we seek an estimate of

$$R(D \to Y) = P(y_1|set(d_1) - P(y_1|set(d_0))$$
  
= 
$$\sum_{u} [P(y_1|d_1, u) - P(y_1|d_0, u)]P(u)$$
 (22)

given the observed probabilities  $P(y, d|z_0)$  and  $P(y, d|z_1)$ .

A few algebraic manipulations of (22) (see Appendix) yields an alternative expression for  $R(D \to Y)$ 

$$R(D \to Y) = E\left[\frac{P(y_1|z_1, u) - P(y_1|z_0, u)}{P(d_1|z_1, u) - P(d_1|z_0, u)}\right]$$
(23)

where E stands for the expectation taken over u.

If we think of u as an index characterizing the experimental units (i.e., the subjects) the result is simple and intuitive. It says that for each individual unit u, the indirect causal effect along the chain  $Z \to D \to Y$  is equal to the product of the individual causal effects along the two links of the chain. If all units were to exhibit the same difference in compliance probabilities,  $P(d_1|z_1,u) - P(d_1|z_0,u)$ , we would have the celebrated instrumental variable formula

$$R(D \to Y) = \frac{R(Z \to Y)}{R(Z \to D)} \tag{24}$$

which says that the causal effect  $R(Z \to Y)$  associated with the intent-to-treat needs to be adjusted upward, through division by the partial compliance  $R(Z \to D)$ . This ratio formula is indeed valid in linear regression models and was derived by econometricians as far back as 1940 [Angrist et al. 1993]. In general, however, since the

quantities on the r.h.s. of Eq. (23) cannot be observed directly (only in expectation), the expression for R can become as low as zero and even negative. Still, when an almost-perfect compliance is observed, the unknown quantities P(y|d,u), P(d|z,u), and P(u) do not have the freedom to render  $R(D \to Y)$  substantially different from  $R(Z \to Y)$ , and meaningful bounds can then be obtained on the actual causal effect of the treatment.

The analysis presented in the Appendix yields the following bounds for the two terms on the r.h.s. of (22):

$$\max[P(y_1, d_1|z_1); \ P(y_1, d_1|z_0)] \le P(y_1|set(d_1)) \le 1 - \max[P(y_0, d_1|z_0); \ P(y_0, d_1|z_1)]$$

$$\max[P(y_1, d_0|z_0); \ P(y_1, d_0|z_1)] \le P(y_1|set(d_0)) \le 1 - \max[P(y_0, d_0|z_0); \ P(y_0, d_0|z_1)]$$

$$(26)$$

Choosing appropriate terms to bound the difference  $P(y_1|set(d_1)) - P(y_1|set(d_0))$ , we obtain a lower bound on the causal effect of D on Y:

$$R(D \to Y) \ge R(Z \to Y) - P(y_1, d_0|z_1) - P(y_0, d_1|z_0)$$
(27)

This bound guarantees that the difference between the causal effect of the intent-totreat and the causal effect of the actual treatment could never exceed the sum of two measurable quantities,  $P(y_1, d_0|z_1) + P(y_0, d_1|z_0)$ . While the bounds in Eqs. (25) and (26) are sharp, the one in Eq. (27) can be further improved using linear programming [Balke & Pearl 1993], albeit at the expense of formal elegance.

Before continuing to Rubin's approach, we should mention that the structural model of Figure 5 imposes definite constraints, obtained directly from Eq. (25)-(26), on the observed distributions  $P(y, d|z_0)$  and  $P(y, d|z_1)$ :

$$P(y_{1}, d_{1}|z_{1}) \leq 1 - P(y_{0}, d_{1}|z_{0})$$

$$P(y_{1}, d_{1}|z_{0}) \leq 1 - P(y_{0}, d_{1}|z_{1})$$

$$P(y_{1}, d_{0}|z_{1}) \leq 1 - P(y_{0}, d_{0}|z_{0})$$

$$P(y_{1}, d_{0}|z_{0}) \leq 1 - P(y_{0}, d_{0}|z_{1})$$
(28)

These constraints constitute necessary and sufficient conditions for a marginal probability P(y,d,z) to be generated by the structure of the model given in Figure 5 and therefore may serve as an operational test for the consistency of that structure with the observed data.

### 7.3 The counterfactual approach

A peculiar feature of the graphical model discussed so far is its capacity for producing meaningful results while keeping the latent variable U totally unspecified. U may be finite or unbounded, discrete or continuous, ordered or unstructured. Although this generality has the advantage of freeing the analyst from commitment to a particular parametric model, it may turn into an inconvenience when finer mathematical details, such as tighter bounds or maximum likelihood estimates, are needed.

The structure of Figure 6 is similar to that of Figure 5, with the difference that the latent variables R R' have only four states each. We will now show that every model that fits into the general latent structure of Figure 5 can also fit into the finite-variable structure of Figure 6 and, moreover, that the states of the variables R and R' correspond precisely to the components of the causal-effect vector in Rubin's model.

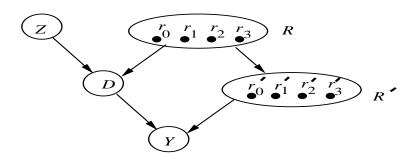


Figure 6: A structure equivalent to that of Figure 5 employing two latent variables, R and R', with 4-states each.

Our first step is to convert each conditional probability term  $P(x_i|\mathbf{pa}_i)$  in Eq. (20) into an equivalent functional form,  $x_i = f_i(\mathbf{pa}_i, \epsilon_i)$ , as in Eq. (6). This can be accomplished by the standard method of simulating probability distributions, letting  $\epsilon_i$  be uniformly distributed over [0,1], and defining

$$X_i = f_i(\mathbf{pa}_i, \epsilon_i) = \begin{cases} 1 & \text{if } \epsilon_i \le P(x_i = 1 | \mathbf{pa}_i) \\ 0 & \text{otherwise} \end{cases}$$
 (29)

It is clear that  $f_i$  induces the specified conditional probability relation between  $\mathbf{pa}_i$  and  $X_i$ .

The next step is to convert the functional equations involving the hypothetical variable U to ones involving variables with a finite number of states. Consider the conditional probability P(d|z,u) in its functional form  $d=f_D(z,u,\epsilon_D)$ . No matter how complex U and  $\epsilon_D$  might be, their impact on D cannot amount to more than a modification of the functional relationship between D and Z and, since there are exactly four functions relating two binary variables, each  $(u,\epsilon_D)$  pair selects one of the four functions. Thus, the impact of the random pair  $(u,\epsilon_D)$  can be simulated by a four-state variable  $r \in \{r_0, r_1, r_2, r_3\}$ , together with the appropriate distribution over r's states, with each state selecting one of the four binary functions.

Formally, if dom(U) and  $dom(\epsilon_D)$  are the domains of U and  $\epsilon_D$ , respectively, define the mapping  $R: dom(U) \times dom(\epsilon_D) \to \{r_0, r_1, r_2, r_3\}$  as follows:

$$R = \begin{cases} r_0 & \text{if } f_D(z_0, u, \epsilon_D) = 0 & \text{and } f_D(z_1, u, \epsilon_D) = 0 \\ r_1 & \text{if } f_D(z_0, u, \epsilon_D) = 0 & \text{and } f_D(z_1, u, \epsilon_D) = 1 \\ r_2 & \text{if } f_D(z_0, u, \epsilon_D) = 1 & \text{and } f_D(z_1, u, \epsilon_D) = 0 \\ r_3 & \text{if } f_D(z_0, u, \epsilon_D) = 1 & \text{and } f_D(z_1, u, \epsilon_D) = 1 \end{cases}$$
(30)

We can now write D as a function of the variables Z and R:

$$d = F_D(z, r) = \begin{cases} d_0 & \text{if} \quad r = r_0 \\ d_0 & \text{if} \quad r = r_1 \quad z = z_0 \\ d_1 & \text{if} \quad r = r_1 \quad z = z_1 \end{cases}$$

$$d_1 & \text{if} \quad r = r_2 \quad z = z_0$$

$$d_0 & \text{if} \quad r = r_2 \quad z = z_1$$

$$d_1 & \text{if} \quad r = r_3$$

$$(31)$$

Repeating the same transformation on the factor P(y|d,u) or its functional form  $y = f_Y(d,u,\epsilon_Y)$  permits us to express Y as a function of D and a second four-state variable R':

$$y = F_Y(d, r') = \begin{cases} y_0 & \text{if } r' = r'_0 \\ y_0 & \text{if } r' = r'_1 & d = d_0 \\ y_1 & \text{if } r' = r'_1 & d = d_1 \\ y_1 & \text{if } r' = r'_2 & d = d_0 \\ y_0 & \text{if } r' = r'_2 & d = d_1 \end{cases}$$

$$(32)$$

$$y_1 & \text{if } r' = r'_2 & d = d_1 \\ y_1 & \text{if } r' = r'_3 & d = d_1 \end{cases}$$

where  $R': dom(U) \times dom(\epsilon_Y) \to \{r'_0, r'_1, r'_2, r'_3\}$ . Since U influences both R and R', the two variables are not independent, hence the arrow  $R \to R'$  in Figure 6. The join distribution over  $R \times R'$  requires 15 independent parameters, and these parameters are sufficient for specifying the model of Figure 6, since Y and D stand in a functional relation to their parents.

The correspondence between the states of variables R and R' and the causal effect vectors in Rubin's model is rather transparent: each state corresponds to a functional relationship between two observed variables. The transformations shown in Eqs. (29) - (32) demonstrate that the so-called counterfactual events emerge in a natural way from a purely mathematical exercise aimed at reducing the domain of the latent variables to the bare minimum.

The causal effect of the treatment can now be obtained directly from Eq. (32), giving

$$P(y_1|set(d_1)) = P(r' = r'_1) + P(r' = r'_3)$$
  

$$P(y_1|set(d_0)) = P(r' = r'_2) + P(r' = r'_3)$$
(33)

and

$$R(D \to Y) = P(r' = r_1') - P(r' = r_2') \tag{34}$$

The computational advantage of this scheme is two-fold. First, lower bounds on  $R(D \to Y)$  can now be produced by minimizing a linear function over a 15-dimensional vector space, rather than by dealing with the unspecified domain of

U. Second, the constraints that the data  $P(y, d|z_0)$  and  $P(y, d|z_1)$  induce on the parameters of P(r, r') are linear, while the constraints induced on the parameters P(d|z, u) and P(y|d, u) in the previous model are non-convex (see Eq. (42) in the Appendix). These advantages enables the use of linear programming techniques to obtain tighter bounds on the causal effect  $R(D \to Y)$  [Balke & Pearl 1993]; such bounds are much harder to obtain in a model where U remains unspecified.

## 8 Conclusions

I hope this paper convinces the reader that DAGs can be used not only for specifying assumptions of conditional independence but also as a formal language for organizing claims about external interventions and their interactions. I hope to have demonstrated as well that DAGs can serve as an analytical tool for quantifying, from non-experimental data, the effect of actions (given qualitative causal structure), for specifying and testing conditions under which randomized experiments are not necessary, and for aiding experimental design and model selection.

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## Appendix

This appendix contains derivations of Eqs. (23) and (27).

To prove (23), we use the conditional independence assumption of (19), and write

$$P(y|z,u) = \sum_{d} P(y|z,d,u) P(d|z,u)$$
 (35)

$$= \sum_{d} P(y|d,u) P(d|z,u)$$
 (36)

which amounts to two equations,

$$P(y_1|z_1,u) = P(y_1|d_1,u) P(d_1|z_1,u) + P(y_1|d_0,u)[1 - P(d_1|z_1,u)]$$

$$P(y_1|z_0,u) = P(y_1|d_1,u) P(d_1|z_0,u) + P(y_1|d_0,u)[1 - P(d_1|z_0,u)]$$
(37)

Solving for  $P(y_1|d_1,u)$  and  $P(y_1|d_0,u)$ , and taking their difference, gives

$$P(y_1|d_1, u) - P(y_1|d_0, u) = \frac{P(y_1|z_1, u) - P(y_1|z_0, u)}{P(d_1|z_1, u) - P(d_1|z_0, u)}$$
(38)

Finally, taking the expectation (over u) on both sides, gives Eq. (23).

To prove (27), we write

$$P(y, d|z) = \sum_{u} P(y|d, u) P(d|z, u) P(u)$$
(39)

and define the following four functions:

$$f_0(u) = P(y_1|d_0, u)$$
  $g_0(u) = P(d_1|u, z_0)$  (40)

$$f_1(u) = P(y_1|d_1, u)$$
  $g_1(u) = P(d_1|u, z_1)$  (41)

This permits us to express six independent components of P(y, d|z) as expectations of these functions:

$$P(y_{1}, d_{0}|z_{0}) = E[f_{0}(1 - g_{0})] = a$$

$$P(y_{1}, d_{0}|z_{1}) = E[f_{0}(1 - g_{1})] = b$$

$$P(d_{1}|z_{0}) = E(g_{0}) = c$$

$$P(d_{1}|z_{1}) = E(g_{1}) = d$$

$$P(y_{1}, d_{1}|z_{0}) = E[f_{1} \cdot g_{0}] = e$$

$$P(y_{1}, d_{1}|z_{1}) = E[f_{1} \cdot g_{1}] = h$$

$$(42)$$

For any two random variables X and Y such that  $0 \le X \le 1, 0 \le Y \le 1$  we have

$$1 + E(XY) - E(Y) \ge E(X) \ge E(XY) \tag{43}$$

since  $E[(1-X)(1-Y)] \ge 0$ . This inequality holds for any pair of f, g functions (since they lie between 0 and 1) and we can write:

$$1 + E(f_1g_0) - E(g_0) \geq E(f_1) \geq E(f_1g_0) 
1 + E(f_1g_1) - E(g_1) \geq E(f_1) \geq E(f_1g_1) 
1 + E[f_0(1 - g_0)] - E(1 - g_0) \geq E(f_0) \geq E[f_0(1 - g_0)] 
1 + E[f_0(1 - g_1)] - E(1 - g_1) \geq E(f_0) \geq E[f_0(1 - g_1)]$$
(44)

$$\max[h; e] \leq E(f_1) \leq \min[(1 + e - c); (1 + h - d)]$$
  
$$\max[a; b] \leq E(f_0) \leq \min[(a + c); (b + d)]$$
(45)

Substituting back the P(y, d|z) expressions from (40) and (41), gives Eqs. (25) and (26). Finally, lower bounding  $E(f_1)$  and upper bounding  $E(f_0)$  provides a lower bound for their difference

$$E(f_1) - E(f_0) \ge \max[e; h] - \min[(a+c); (b+d)]$$
  
  $\ge h - (a+c)$  (46)

from which Eq. (27) follows.