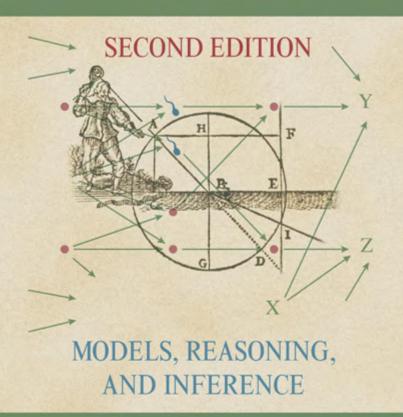
# CAUSALITY



# JUDEA PEARL

## **CAUSALITY**

### Models, Reasoning, and Inference Second Edition

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#### **Preface to the First Edition**

The central aim of many studies in the physical, behavioral, social, and biological sciences is the elucidation of cause–effect relationships among variables or events. However, the appropriate methodology for extracting such relationships from data – or even from theories – has been fiercely debated.

The two fundamental questions of causality are: (1) What empirical evidence is required for legitimate inference of cause–effect relationships? (2) Given that we are willing to accept causal information about a phenomenon, what inferences can we draw from such information, and how? These questions have been without satisfactory answers in part because we have not had a clear semantics for causal claims and in part because we have not had effective mathematical tools for casting causal questions or deriving causal answers.

In the last decade, owing partly to advances in graphical models, causality has undergone a major transformation: from a concept shrouded in mystery into a mathematical object with well-defined semantics and well-founded logic. Paradoxes and controversies have been resolved, slippery concepts have been explicated, and practical problems relying on causal information that long were regarded as either metaphysical or unmanageable can now be solved using elementary mathematics. Put simply, causality has been mathematized.

This book provides a systematic account of this causal transformation, addressed primarily to readers in the fields of statistics, artificial intelligence, philosophy, cognitive science, and the health and social sciences. Following a description of the conceptual and mathematical advances in causal inference, the book emphasizes practical methods for elucidating potentially causal relationships from data, deriving causal relationships from combinations of knowledge and data, predicting the effects of actions and policies, evaluating explanations for observed events and scenarios, and – more generally – identifying and explicating the assumptions needed for substantiating causal claims.

Ten years ago, when I began writing *Probabilistic Reasoning in Intelligent Systems* (1988), I was working within the empiricist tradition. In this tradition, probabilistic relationships constitute the foundations of human knowledge, whereas causality simply provides useful ways of abbreviating and organizing intricate patterns of probabilistic relationships. Today, my view is quite different. I now take causal relationships to be the

fundamental building blocks both of physical reality and of human understanding of that reality, and I regard probabilistic relationships as but the surface phenomena of the causal machinery that underlies and propels our understanding of the world.

Accordingly, I see no greater impediment to scientific progress than the prevailing practice of focusing all of our mathematical resources on probabilistic and statistical inferences while leaving causal considerations to the mercy of intuition and good judgment. Thus I have tried in this book to present mathematical tools that handle causal relationships side by side with probabilistic relationships. The prerequisites are startlingly simple, the results embarrassingly straightforward. No more than basic skills in probability theory and some familiarity with graphs are needed for the reader to begin solving causal problems that are too complex for the unaided intellect. Using simple extensions of probability calculus, the reader will be able to determine mathematically what effects an intervention might have, what measurements are appropriate for control of confounding, how to exploit measurements that lie on the causal pathways, how to trade one set of measurements for another, and how to estimate the probability that one event was the actual cause of another.

Expert knowledge of logic and probability is nowhere assumed in this book, but some general knowledge in these areas is beneficial. Thus, Chapter 1 includes a summary of the elementary background in probability theory and graph notation needed for the understanding of this book, together with an outline of the developments of the last decade in graphical models and causal diagrams. This chapter describes the basic paradigms, defines the major problems, and points readers to the chapters that provide solutions to those problems.

Subsequent chapters include introductions that serve both to orient the reader and to facilitate skipping; they indicate safe detours around mathematically advanced topics, specific applications, and other explorations of interest primarily to the specialist.

The sequence of discussion follows more or less the chronological order by which our team at UCLA has tackled these topics, thus re-creating for the reader some of our excitement that accompanied these developments. Following the introductory chapter (Chapter 1), we start with the hardest questions of how one can go about discovering cause-effect relationships in raw data (Chapter 2) and what guarantees one can give to ensure the validity of the relationships thus discovered. We then proceed to questions of identifiability - namely, predicting the direct and indirect effects of actions and policies from a combination of data and fragmentary knowledge of where causal relationships might operate (Chapters 3 and 4). The implications of these findings for the social and health sciences are then discussed in Chapters 5 and 6 (respectively), where we examine the concepts of structural equations and confounding. Chapter 7 offers a formal theory of counterfactuals and structural models, followed by a discussion and a unification of related approaches in philosophy, statistics, and economics. The applications of counterfactual analysis are then pursued in Chapters 8–10, where we develop methods of bounding causal relationships and illustrate applications to imperfect experiments, legal responsibility, and the probability of necessary, sufficient, and single-event causation. We end this book (Epilogue) with a transcript of a public lecture that I presented at UCLA, which provides a gentle introduction to the historical and conceptual aspects of causation.

Readers who wish to be first introduced to the nonmathematical aspects of causation are advised to start with the Epilogue and then to sweep through the other historical/conceptual parts of the book: Sections 1.1.1, 3.3.3, 4.5.3, 5.1, 5.4.1, 6.1, 7.2, 7.4, 7.5, 8.3, 9.1, 9.3, and 10.1. More formally driven readers, who may be anxious to delve directly into the mathematical aspects and computational tools, are advised to start with Section 7.1 and then to proceed as follows for tool building: Section 1.2, Chapter 3, Sections 4.2–4.4, Sections 5.2–5.3, Sections 6.2–6.3, Section 7.3, and Chapters 8–10.

I owe a great debt to many people who assisted me with this work. First, I would like to thank the members of the Cognitive Systems Laboratory at UCLA, whose work and ideas formed the basis of many of these sections: Alex Balke, Blai Bonet, David Chickering, Adnan Darwiche, Rina Dechter, David Galles, Hector Geffner, Dan Geiger, Moisés Goldszmidt, Jin Kim, Jin Tian, and Thomas Verma. Tom and Dan have proven some of the most basic theorems in causal graphs; Hector, Adnan, and Moisés were responsible for keeping me in line with the logicist approach to actions and change; and Alex and David have taught me that counterfactuals are simpler than the name may imply.

My academic and professional colleagues have been very generous with their time and ideas as I began ploughing the peaceful territories of statistics, economics, epidemiology, philosophy, and the social sciences. My mentors—listeners in statistics have been Phil Dawid, Steffen Lauritzen, Don Rubin, Art Dempster, David Freedman, and David Cox. In economics, I have benefited from many discussions with John Aldrich, Kevin Hoover, James Heckman, Ed Learner, and Herbert Simon. My forays into epidemiology resulted in a most fortunate and productive collaboration with Sander Greenland and James Robins. Philosophical debates with James Woodward, Nancy Cartwright, Brian Skyrms, Clark Glymour, and Peter Spirtes have sharpened my thinking of causality in and outside philosophy. Finally, in artificial intelligence, I have benefited from discussions with and the encouragement of Nils Nilsson, Ray Reiter, Don Michie, Joe Halpern, and David Heckerman.

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J. P. Los Angeles August 1999

#### Preface to the Second Edition

It has been more than eight years since the first edition of this book presented readers with the friendly face of causation and her mathematical artistry. The popular reception of the book and the rapid expansion of the structural theory of causation call for a new edition to assist causation through her second transformation – from a demystified wonder to a commonplace tool in research and education. This edition (1) provides technical corrections, updates, and clarifications in all ten chapters of the original book, (2) adds summaries of new developments and annotated bibliographical references at the end of each chapter, and (3) elucidates subtle issues that readers and reviewers have found perplexing, objectionable, or in need of elaboration. These are assembled into an entirely new chapter (11) which, I sincerely hope, clears the province of causal thinking from the last traces of controversy.

Teachers who have taught from this book before should find the revised edition more lucid and palatable, while those who have waited for scouts to carve the path will find the road paved and tested. Supplementary educational material, slides, tutorials, and homework can be found on my website, http://www.cs.ucla.edu/~judea/.

My main audience remain the students: students of statistics who wonder why instructors are reluctant to discuss causality in class; students of epidemiology who wonder why elementary concepts such as confounding are so hard to define mathematically; students of economics and social science who question the meaning of the parameters they estimate; and, naturally, students of artificial intelligence and cognitive science, who write programs and theories for knowledge discovery, causal explanations, and causal speech.

I hope that each of these groups will find the unified theory of causation presented in this book to be both inspirational and instrumental in tackling new challenges in their respective fields.

J. P. Los Angeles July 2008

## Introduction to Probabilities, Graphs, and Causal Models

Chance gives rise to thoughts, and chance removes them. Pascal (1670)

#### 1.1 INTRODUCTION TO PROBABILITY THEORY

#### 1.1.1 Why Probabilities?

Causality connotes lawlike necessity, whereas probabilities connote exceptionality, doubt, and lack of regularity. Still, there are two compelling reasons for starting with, and in fact stressing, probabilistic analysis of causality; one is fairly straightforward, the other more subtle.

The simple reason rests on the observation that causal utterances are often used in situations that are plagued with uncertainty. We say, for example, "reckless driving causes accidents" or "you will fail the course because of your laziness" (Suppes 1970), knowing quite well that the antecedents merely tend to make the consequences more likely, not absolutely certain. Any theory of causality that aims at accommodating such utterances must therefore be cast in a language that distinguishes various shades of likelihood – namely, the language of probabilities. Connected with this observation, we note that probability theory is currently the official mathematical language of most disciplines that use causal modeling, including economics, epidemiology, sociology, and psychology. In these disciplines, investigators are concerned not merely with the presence or absence of causal connections but also with the relative strengths of those connections and with ways of inferring those connections from noisy observations. Probability theory, aided by methods of statistical analysis, provides both the principles and the means of coping with – and drawing inferences from – such observations.

The more subtle reason concerns the fact that even the most assertive causal expressions in natural language are subject to exceptions, and those exceptions may cause major difficulties if processed by standard rules of deterministic logic. Consider, for example, the two plausible premises:

- 1. My neighbor's roof gets wet whenever mine does.
- 2. If I hose my roof it will get wet.

Taken literally, these two premises imply the implausible conclusion that my neighbor's roof gets wet whenever I hose mine.

Such paradoxical conclusions are normally attributed to the finite granularity of our language, as manifested in the many exceptions that are implicit in premise 1. Indeed, the paradox disappears once we take the trouble of explicating those exceptions and write, for instance:

1\*. My neighbor's roof gets wet whenever mine does, except when it is covered with plastic, or when my roof is hosed, etc.

Probability theory, by virtue of being especially equipped to tolerate unexplicated exceptions, allows us to focus on the main issues of causality without having to cope with paradoxes of this kind.

As we shall see in subsequent chapters, tolerating exceptions solves only some of the problems associated with causality. The remaining problems – including issues of inference, interventions, identification, ramification, confounding, counterfactuals, and explanation – will be the main topic of this book. By portraying those problems in the language of probabilities, we emphasize their universality across languages. Chapter 7 will recast these problems in the language of deterministic logic and will introduce probabilities merely as a way to express uncertainty about unobserved facts.

#### 1.1.2 Basic Concepts in Probability Theory

The bulk of the discussion in this book will focus on systems with a finite number of discrete variables and thus will require only rudimentary notation and elementary concepts in probability theory. Extensions to continuous variables will be outlined but not elaborated in full generality. Readers who want additional mathematical machinery are invited to study the many excellent textbooks on the subject – for example, Feller (1950), Hoel et al. (1971), or the appendix to Suppes (1970). This section provides a brief summary of elementary probability concepts, based largely on Pearl (1988b), with special emphasis on Bayesian inference and its connection to the psychology of human reasoning under uncertainty. Such emphasis is generally missing from standard textbooks.

We will adhere to the Bayesian interpretation of probability, according to which probabilities encode degrees of belief about events in the world and data are used to strengthen, update, or weaken those degrees of belief. In this formalism, degrees of belief are assigned to propositions (sentences that take on true or false values) in some language, and those degrees of belief are combined and manipulated according to the rules of probability calculus. We will make no distinction between sentential propositions and the actual events represented by those propositions. For example, if A stands for the statement "Ted Kennedy will seek the nomination for president in year 2012," then  $P(A \mid K)$  stands for a person's subjective belief in the event described by A given a body of knowledge K, which might include that person's assumptions about American politics, specific proclamations made by Kennedy, and an assessment of Kennedy's age and personality. In defining probability expressions, we often simply write P(A), leaving out the symbol K. However, when the background information undergoes changes, we need to identify specifically the assumptions that account for our beliefs and explicitly articulate K (or some of its elements).

In the Bayesian formalism, belief measures obey the three basic axioms of probability calculus:

$$0 \le P(A) \le 1,\tag{1.1}$$

$$P$$
 (sure proposition) = 1, (1.2)

$$P(A \text{ or } B) = P(A) + P(B) \text{ if } A \text{ and } B \text{ are mutually exclusive.}$$
 (1.3)

The third axiom states that the belief assigned to any set of events is the sum of the beliefs assigned to its nonintersecting components. Because any event A can be written as the union of the joint events  $(A \wedge B)$  and  $(A \wedge \neg B)$ , their associated probabilities are given by A

$$P(A) = P(A, B) + P(A, \neg B),$$
 (1.4)

where P(A, B) is short for  $P(A \land B)$ . More generally, if  $B_i$ , i = 1, 2, ..., n, is a set of exhaustive and mutually exclusive propositions (called a *partition* or a *variable*), then P(A) can be computed from  $P(A, B_i)$ , i = 1, 2, ..., n, by using the sum

$$P(A) = \sum_{i} P(A, B_i),$$
 (1.5)

which has come to be known as the "law of *total* probability." The operation of summing up probabilities over all  $B_i$  is also called "marginalizing over B"; and the resulting probability, P(A), is called the *marginal* probability of A. For example, the probability of A, "The outcomes of two dice are equal," can be computed by summing over the joint events  $(A \land B_i)$ , i = 1, 2, ..., 6, where  $B_i$  stands for the proposition "The outcome of the first die is i." This yields

$$P(A) = \sum_{i} P(A, B_i) = 6 \times \frac{1}{36} = \frac{1}{6}.$$
 (1.6)

A direct consequence of (1.2) and (1.4) is that a proposition and its negation must be assigned a total belief of unity,

$$P(A) + P(\neg A) = 1,$$
 (1.7)

because one of the two statements is certain to be true.

The basic expressions in the Bayesian formalism are statements about *conditional* probabilities – for example,  $P(A \mid B)$  – which specify the belief in A under the assumption that B is known with absolute certainty. If  $P(A \mid B) = P(A)$ , we say that A and B are independent, since our belief in A remains unchanged upon learning the truth of B. If  $P(A \mid B, C) = P(A \mid C)$ , we say that A and B are conditionally independent given C; that is, once we know C, learning B would not change our belief in A.

Contrary to the traditional practice of defining conditional probabilities in terms of joint events,

$$P(A \mid B) = \frac{P(A, B)}{P(B)},$$
 (1.8)

<sup>&</sup>lt;sup>1</sup> The symbols  $\land$ ,  $\lor$ ,  $\neg$ ,  $\Rightarrow$  denote the logical connectives *and*, *or*, *not*, and *implies*, respectively.

Bayesian philosophers see the conditional relationship as more basic than that of joint events – that is, more compatible with the organization of human knowledge. In this view, B serves as a pointer to a context or frame of knowledge, and  $A \mid B$  stands for an event A in the context specified by B (e.g., a symptom A in the context of a disease B). Consequently, empirical knowledge invariably will be encoded in conditional probability statements, whereas belief in joint events (if it is ever needed) will be computed from those statements via the product

$$P(A,B) = P(A|B)P(B), \tag{1.9}$$

which is equivalent to (1.8). For example, it was somewhat unnatural to assess

$$P(A, B_i) = \frac{1}{36}$$

directly in (1.6). The mental process underlying such assessment presumes that the two outcomes are independent, so to make this assumption explicit the probability of the joint event (equality,  $B_i$ ) should be assessed from the conditional event (equality  $|B_i\rangle$ ) via the product

$$P(\text{equality } \mid B_i) P(B_i) = P(\text{outcome of second die is } i \mid B_i) P(B_i)$$
$$= \frac{1}{6} \times \frac{1}{6} = \frac{1}{36}.$$

As in (1.5), the probability of any event A can be computed by conditioning it on any set of exhaustive and mutually exclusive events  $B_i$ , i = 1, 2, ..., n, and then summing:

$$P(A) = \sum_{i} P(A \mid B_{i}) P(B_{i}). \tag{1.10}$$

This decomposition provides the basis for hypothetical or "assumption-based" reasoning. It states that the belief in any event A is a weighted sum over the beliefs in all the distinct ways that A might be realized. For example, if we wish to calculate the probability that the outcome X of the first die will be greater than the outcome Y of the second, we can condition the event A: X > Y on all possible values of X and obtain

$$P(A) = \sum_{i=1}^{6} P(Y < X | X = i) P(X = i)$$

$$= \sum_{i=1}^{6} P(Y < i) \frac{1}{6} = \sum_{i=1}^{6} \sum_{j=1}^{i-1} P(Y = j) \frac{1}{6}$$

$$= \frac{1}{6} \sum_{i=2}^{6} \frac{i-1}{6} = \frac{5}{12}.$$

It is worth reemphasizing that formulas like (1.10) are always understood to apply in some larger context K, which defines the assumptions taken as common knowledge (e.g., the fairness of dice rolling). Equation (1.10) is really a shorthand notation for the statement

$$P(A \mid K) = \sum_{i} P(A \mid B_{i}, K) P(B_{i} \mid K).$$
(1.11)

This equation follows from the fact that every conditional probability  $P(A \mid K)$  is itself a genuine probability function; hence it satisfies (1.10).

Another useful generalization of the product rule (equation (1.9)) is the *chain rule* formula. It states that if we have a set of n events,  $E_1, E_2, \ldots, E_n$ , then the probability of the joint event  $(E_1, E_2, \ldots, E_n)$  can be written as a product of n conditional probabilities:

$$P(E_1, E_2, \dots, E_n) = P(E_n \mid E_{n-1}, \dots, E_2, E_1) \dots P(E_2 \mid E_1) P(E_1).$$
(1.12)

This product can be derived by repeated application of (1.9) in any convenient order. The heart of Bayesian inference lies in the celebrated inversion formula,

$$P(H \mid e) = \frac{P(e \mid H)P(H)}{P(e)},\tag{1.13}$$

which states that the belief we accord a hypothesis H upon obtaining evidence e can be computed by multiplying our previous belief P(H) by the likelihood  $P(e \mid H)$  that e will materialize if H is true. This  $P(H \mid e)$  is sometimes called the posterior probability (or simply *posterior*), and P(H) is called the prior probability (or *prior*). The denominator P(e) of (1.13) hardly enters into consideration because it is merely a normalizing constant  $P(e) = P(e \mid H)P(H) + P(e \mid \neg H)P(\neg H)$ , which can be computed by requiring that  $P(H \mid e)$  and  $P(\neg H \mid e)$  sum to unity.

Whereas formally (1.13) might be dismissed as a tautology stemming from the definition of conditional probabilities,

$$P(A \mid B) = \frac{P(A, B)}{P(B)}$$
 and  $P(B \mid A) = \frac{P(A, B)}{P(A)}$ , (1.14)

the Bayesian subjectivist regards (1.13) as a normative rule for updating beliefs in response to evidence. In other words, although conditional probabilities can be viewed as purely mathematical constructs (as in (1.14)), the Bayes adherent views them as primitives of the language and as faithful translations of the English expression "…, given that I know A." Accordingly, (1.14) is not a definition but rather an empirically verifiable relationship between English expressions. It asserts, among other things, that the belief a person attributes to B after discovering A is never lower than that attributed to  $A \wedge B$  before discovering A. Also, the ratio between these two beliefs will increase proportionally with the degree of surprise  $[P(A)]^{-1}$  one associates with the discovery of A.

The importance of (1.13) is that it expresses a quantity  $P(H \mid e)$  – which people often find hard to assess – in terms of quantities that often can be drawn directly from our experiential knowledge. For example, if a person at the next gambling table declares the outcome "twelve," and we wish to know whether he was rolling a pair of dice or spinning a roulette wheel, our models of the gambling devices readily yield the quantities  $P(\text{twelve} \mid \text{dice})$  and  $P(\text{twelve} \mid \text{roulette})$ : 1/36 for the former and 1/38 for the latter. Similarly, we can judge the prior probabilities P(dice) and P(roulette) by estimating the number of roulette wheels and dice tables at the casino. Issuing a direct judgment of

 $P(\text{dice} \mid \text{twelve})$  would have been much more difficult; only a specialist in such judgments, trained at the very same casino, could do it reliably.

In order to complete this brief introduction, we must discuss the notion of *probabilistic model* (also called *probability space*). A probabilistic model is an encoding of information that permits us to compute the probability of every well-formed sentence S in accordance with the axioms of (1.1)–(1.3). Starting with a set of atomic propositions  $A, B, C, \ldots$ , the set of well-formed sentences consists of all Boolean formulas involving these propositions, for example,  $S = (A \land B) \lor \neg C$ . The traditional method of specifying probabilistic models employs a *joint distribution function*, which is a function that assigns nonnegative weights to every *elementary event* in the language (an elementary event being a conjunction in which every atomic proposition or its negation appears once) such that the sum of the weights adds up to 1. For example, if we have three atomic propositions, A, B, and C, then a joint distribution function should assign nonnegative weights to all eight combinations  $-(A \land B \land C), (A \land B \neg C), \ldots, (\neg A \land \neg B \land \neg C)$  – such that the eight weights sum to 1.

The reader may recognize the set of elementary events as the *sample space* in probability textbooks. For example, if A, B, and C correspond to the propositions that coins 1, 2, and 3 will come up heads, then the sample space will consist of the set {HHH, HHT, HTH,..., TTT}. Indeed, it is sometimes convenient to view the conjunctive formulas corresponding to elementary events as *points* (or *worlds* or *configurations*), and to regard other formulas as *sets* made up of these points. Since every Boolean formula can be expressed as a disjunction of elementary events, and since the elementary events are mutually exclusive, we can always compute P(S) using the additivity axiom (equation (1.3)). Conditional probabilities can be computed the same way, using (1.14). Thus, any joint probability function represents a complete probabilistic model.

Joint distribution functions are mathematical constructs of great importance. They allow us to determine quickly whether we have sufficient information to specify a complete probabilistic model, whether the information we have is consistent, and at what point additional information is needed. The criteria are simply to check (i) whether the information available is sufficient for uniquely determining the probability of every elementary event in the domain and (ii) whether the probabilities add up to 1.

In practice, however, joint distribution functions are rarely specified explicitly. In the analysis of continuous random variables, the distribution functions are given by algebraic expressions such as those describing normal or exponential distributions; for discrete variables, indirect representation methods have been developed where the overall distribution is inferred from local relationships among small groups of variables. Graphical models, the most popular of these representations, provide the basis of discussion throughout this book. Their use and formal characterization will be discussed in the next few sections.

#### 1.1.3 Combining Predictive and Diagnostic Supports

The essence of Bayes's rule (equation 1.13)) is conveniently portrayed using the *odds* and *likelihood ratio* parameters. Dividing (1.13) by the complementary form for  $P(\neg H \mid e)$ , we obtain

$$\frac{P(H \mid e)}{P(\neg H \mid e)} = \frac{P(e \mid H)}{P(e \mid \neg H)} \frac{P(H)}{P(\neg H)}.$$
(1.15)

Defining the prior odds on H as

$$O(H) = \frac{P(H)}{P(\neg H)} = \frac{P(H)}{1 - P(H)}$$
(1.16)

and the likelihood ratio as

$$L(e \mid H) = \frac{P(e \mid H)}{P(e \mid \neg H)},\tag{1.17}$$

the posterior odds

$$O(H \mid e) = \frac{P(H \mid e)}{P(\neg H \mid e)} \tag{1.18}$$

are given by the product

$$O(H \mid e) = L(e \mid H)O(H).$$
 (1.19)

Thus, Bayes's rule dictates that the overall strength of belief in a hypothesis H, based on both our previous knowledge K and the observed evidence e, should be the product of two factors: the prior odds O(H) and the likelihood ratio  $L(e \mid H)$ . The first factor measures the *predictive* or *prospective* support accorded to H by the background knowledge alone, while the second represents the *diagnostic* or *retrospective* support given to H by the evidence actually observed.<sup>2</sup>

Strictly speaking, the likelihood ratio  $L(e \mid H)$  might depend on the content of the tacit knowledge base K. However, the power of Bayesian techniques comes primarily from the fact that, in causal reasoning, the relationship  $P(e \mid H)$  is fairly local: given that H is true, the probability of e can be estimated naturally since it is usually not dependent on many other propositions in the knowledge base. For example, once we establish that a patient suffers from a given disease H, it is natural to estimate the probability that she will develop a certain symptom e. The organization of medical knowledge rests on the paradigm that a symptom is a stable characteristic of the disease and should therefore be fairly independent of other factors, such as epidemic conditions, previous diseases, and faulty diagnostic equipment. For this reason the conditional probabilities  $P(e \mid H)$ , as opposed to  $P(H \mid e)$ , are the atomic relationships in Bayesian analysis. The former possess modularity features similar to logical rules. They convey a degree of confidence in rules such as "If H then e," a confidence that persists regardless of what other rules or facts reside in the knowledge base.

**Example 1.1.1** Imagine being awakened one night by the shrill sound of your burglar alarm. What is your degree of belief that a burglary attempt has taken place? For

<sup>&</sup>lt;sup>2</sup> In epidemiology, if H stands for exposure and e stands for disease, then the likelihood ratio L is called the "risk ratio" (Rothman and Greenland 1998, p. 50). Equation (1.18) would then give the odds that a person with disease e had been exposed to H.

illustrative purposes we make the following judgments: (a) There is a 95% chance that an attempted burglary will trigger the alarm system  $-P(\text{alarm} \mid \text{burglary}) = 0.95$ ; (b) based on previous false alarms, there is a slight (1%) chance that the alarm will be triggered by a mechanism other than an attempted burglary  $-P(\text{alarm} \mid \text{no burglary}) = 0.01$ ; (c) previous crime patterns indicate that there is a one in ten thousand chance that a given house will be burglarized on a given night  $-P(\text{burglary}) = 10^{-4}$ .

Putting these assumptions together using (1.19), we obtain

 $O(\text{burglary} \mid \text{alarm}) = L(\text{alarm} \mid \text{burglary}) O(\text{burglary})$ 

$$= \frac{0.95}{0.01} \frac{10^{-4}}{1 - 10^{-4}} = 0.0095.$$

So, from

$$P(A) = \frac{O(A)}{1 + O(A)} \tag{1.20}$$

we have

$$P(\text{burglary} \mid \text{alarm}) = \frac{0.0095}{1 + 0.0095} = 0.00941.$$

Thus, the retrospective support imparted to the burglary hypothesis by the alarm evidence has increased its degree of belief almost a hundredfold, from one in ten thousand to 94.1 in ten thousand. The fact that the belief in burglary is still below 1% should not be surprising, given that the system produces a false alarm almost once every three months. Notice that it was not necessary to estimate the absolute values of the probabilities  $P(\text{alarm} \mid \text{burglary})$  and  $P(\text{alarm} \mid \text{no burglary})$ . Only their ratio enters the calculation, so a direct estimate of this ratio could have been used instead.

#### 1.1.4 Random Variables and Expectations

By a *variable* we will mean an attribute, measurement or inquiry that may take on one of several possible outcomes, or *values*, from a specified domain. If we have beliefs (i.e., probabilities) attached to the possible values that a variable may attain, we will call that variable a *random variable*. For example, the color of the shoes that I will wear tomorrow is a random variable named "color," and the values it may take come from the domain {yellow, green, red,...}.

Most of our analysis will concern a finite set V of random variables (also called *partitions*) where each variable  $X \in V$  may take on values from a finite domain  $D_X$ . We will use capital letters (e.g., X, Y, Z) for variable names and lowercase letters (x, y, z)

This is a minor generalization of the textbook definition, according to which a random variable is a mapping from the sample space (e.g., the set of elementary events) to the real line. In our definition, the mapping is from the sample space to any set of objects called "values," which may or may not be ordered.

as generic symbols for specific values taken by the corresponding variables. For example, if X stands for the color of an object, then x will designate any possible choice of an element from the set {yellow, green, red,...}. Clearly, the proposition X = yellow describes an event, namely, a subset of possible states of affair that satisfy the proposition "the color of the object is yellow." Likewise, each variable X can be viewed as a partition of the states of the world, since the statement X = x defines a set of exhaustive and mutually exclusive sets of states, one for each value of x.

In most of our discussions, we will not make notational distinction between variables and sets of variables, because a set of variables essentially defines a compound variable whose domain is the Cartesian product of the domains of the individual constituents in the set. Thus, if Z stands for the set  $\{X, Y\}$ , then z stands for pairs (x, y) such that  $x \in D_X$  and  $y \in D_Y$ . When the distinction between variables and sets of variables requires special emphasis, indexed letters (say,  $X_1, X_2, ..., X_n$  or  $V_1, V_2, ..., V_n$ ) will be used to represent individual variables.

We shall consistently use the abbreviation P(x) for the probabilities P(X = x),  $x \in D_x$ . Likewise, if Z stands for the set  $\{X, Y\}$ , then P(z) will be defined as

$$P(z) \triangleq P(Z=z) = P(X=x, Y=y), x \in D_X, y \in D_Y.$$

When the values of a random variable *X* are real numbers, *X* is called a *real* random variable; one can then define the *mean* or *expected value* of *X* as

$$E(X) \triangleq \sum_{x} x P(x) \tag{1.21}$$

and the conditional mean of X, given event Y = y, as

$$E(X \mid y) \triangleq \sum_{x} x P(x \mid y). \tag{1.22}$$

The expectation of any function g of X is defined as

$$E[g(X)] \triangleq \sum_{x} g(x)P(x). \tag{1.23}$$

In particular, the function  $g(X) = (X - E(X))^2$  has received much attention; its expectation is called the *variance* of X, denoted  $\sigma_X^2$ ;

$$\sigma_X^2 \triangleq E[(X - E(X))^2].$$

The conditional mean  $E(X \mid Y = y)$  is the *best estimate* of X, given the observation Y = y, in the sense of minimizing the expected square error  $\sum_{x} (x - x')^2 P(x \mid y)$  over all possible x'.

The expectation of a function g(X, Y) of two variables, X and Y, requires the joint probability P(x, y) and is defined as

$$E[g(X, Y)] \triangleq \sum_{x, y} g(x, y) P(x, y)$$

(cf. equation (1.23)). Of special importance is the expectation of the product (g(X, Y) = (X - E(X))(Y - E(Y)), which is known as the *covariance* of X and Y,

$$\sigma_{XY} \triangleq E[(X - E(X))(Y - E(Y))],$$

and which is often normalized to yield the correlation coefficient

$$\rho_{XY} = \frac{\sigma_{XY}}{\sigma_X \sigma_Y}$$

and the regression coefficient (of X on Y)

$$r_{XY} \stackrel{\triangle}{=} \rho_{XY} \frac{\sigma_X}{\sigma_Y} = \frac{\sigma_{XY}}{\sigma_Y^2}.$$

The *conditional* variance, covariance, and correlation coefficient, given Z = z, are defined in a similar manner, using the conditional distribution  $P(x, y \mid z)$  in taking expectations. In particular, the *conditional correlation coefficient*, given Z = z, is defined as

$$\rho_{XY|z} = \frac{\sigma_{XY|z}}{\sigma_{X|z}\,\sigma_{Y|z}}.\tag{1.24}$$

Additional properties, specific to normal distributions, will be reviewed in Chapter 5 (Section 5.2.1).

The foregoing definitions apply to discrete random variables – that is, variables that take on finite or denumerable sets of values on the real line. The treatment of expectation and correlation is more often applied to continuous random variables, which are characterized by a *density function* f(x) defined as follows:

$$P(a \le X \le b) = \int_{a}^{b} f(x) \, dx$$

for any two real numbers a and b with a < b. If X is discrete, then f(x) coincides with the probability function P(x), once we interpret the integral through the translation

$$\int_{-\infty}^{\infty} f(x)dx \iff \sum_{x} P(x). \tag{1.25}$$

Readers accustomed to continuous analysis should bear this translation in mind whenever summation is used in this book. For example, the expected value of a continuous random variable X can be obtained from (1.21), to read

$$E(X) = \int_{-\infty}^{\infty} x f(x) dx,$$

with analogous translations for the variance, correlation, and so forth.

We now turn to define *conditional independence* relationships among variables, a central notion in causal modelling.

#### 1.1.5 Conditional Independence and Graphoids

#### **Definition 1.1.2 (Conditional Independence)**

Let  $V = \{V_1, V_2, ...\}$  be a finite set of variables. Let  $P(\cdot)$  be a joint probability function over the variables in V, and let X, Y, Z stand for any three subsets of variables in V. The sets X and Y are said to be conditionally independent given Z if

$$P(x \mid y, z) = P(x \mid z) \quad \text{whenever} \quad P(y, z) > 0. \tag{1.26}$$

In words, learning the value of Y does not provide additional information about X, once we know Z. (Metaphorically, Z "screens off" X from Y.)

Equation (1.26) is a terse way of saying the following: For any configuration x of the variables in the set X and for any configurations y and z of the variables in Y and Z satisfying P(Y = y, Z = z) > 0, we have

$$P(X = x | Y = y, Z = z) = P(X = x | Z = z).$$
(1.27)

We will use Dawid's (1979) notation  $(X \perp\!\!\!\perp Y \mid Z)_P$  or simply  $(X \perp\!\!\!\perp Y \mid Z)$  to denote the conditional independence of X and Y given Z; thus,

$$(X \perp \!\!\!\perp Y \mid Z)_P \quad \text{iff} \quad P(x \mid y, z) = P(x \mid z) \tag{1.28}$$

for all values x, y, z such that P(y, z) > 0. Unconditional independence (also called marginal independence) will be denoted by  $(X \perp\!\!\!\perp Y \mid \emptyset)$ ; that is,

$$(X \perp \!\!\!\perp Y \mid \emptyset)$$
 iff  $P(x \mid y) = P(x)$  whenever  $P(y) > 0$  (1.29)

("iff" is shorthand for "if and only if"). Note that  $(X \perp\!\!\!\perp Y \mid Z)$  implies the conditional independence of all pairs of variables  $V_i \in X$  and  $V_j \in Y$ , but the converse is not necessarily true.

The following is a (partial) list of properties satisfied by the conditional independence relation  $(X \perp\!\!\!\perp Y \mid Z)$ .

Symmetry:  $(X \perp\!\!\!\perp Y \mid Z) \Longrightarrow (Y \perp\!\!\!\perp X \mid Z)$ .

**Decomposition:**  $(X \perp\!\!\!\perp YW \mid Z) \Longrightarrow (X \perp\!\!\!\perp Y \mid Z)$ .

Weak union:  $(X \perp\!\!\!\perp YW \mid Z) \Longrightarrow (X \perp\!\!\!\perp Y \mid ZW)$ .

Contraction:  $(X \perp\!\!\!\perp Y \mid Z) \& (X \perp\!\!\!\perp W \mid ZY) \Longrightarrow (X \perp\!\!\!\perp YW \mid Z)$ .

*Intersection:*  $(X \perp\!\!\!\perp W \mid ZY) \& (X \perp\!\!\!\perp Y \mid ZW) \Longrightarrow (X \perp\!\!\!\perp YW \mid Z).$ 

(Intersection is valid in strictly positive probability distributions.)

The proof of these properties can be derived by elementary means from (1.28) and the basic axioms of probability theory. <sup>4</sup> These properties were called *graphoid axioms* by

<sup>&</sup>lt;sup>4</sup> These properties were first introduced by Dawid (1979) and Spohn (1980) in a slightly different form, and were independently proposed by Pearl and Paz (1987) to characterize the relationships between graphs and informational relevance. Geiger and Pearl (1993) present an in-depth analysis.

Pearl and Paz (1987) and Geiger et al. (1990) and have been shown to govern the concept of informational relevance in a wide variety of interpretations (Pearl 1988b). In graphs, for example, these properties are satisfied if we interpret  $(X \perp \!\!\! \perp \!\!\! \perp \!\!\! Y \mid Z)$  to mean "all paths from a subset X of nodes to a subset Y of nodes are intercepted by a subset Z of nodes."

The intuitive interpretation of the graphoid axioms is as follows (Pearl 1988b, p. 85). The *symmetry* axiom states that, in any state of knowledge Z, if Y tells us nothing new about X, then X tells us nothing new about Y. The *decomposition* axiom asserts that if two combined items of information are judged irrelevant to X, then each separate item is irrelevant as well. The *weak union* axiom states that learning irrelevant information W cannot help the irrelevant information Y become relevant to X. The *contraction* axiom states that if we judge W irrelevant to X after learning some irrelevant information Y, then W must have been irrelevant before we learned Y. Together, the weak union and contraction properties mean that irrelevant information should not alter the relevance status of other propositions in the system; what was relevant remains relevant, and what was irrelevant remains irrelevant. The *intersection* axiom states that if Y is irrelevant to X when we know W and if W is irrelevant to X when we know Y, then neither W nor Y (nor their combination) is relevant to X.

#### 1.2 GRAPHS AND PROBABILITIES

#### 1.2.1 Graphical Notation and Terminology

A graph consists of a set V of vertices (or nodes) and a set E of edges (or links) that connect some pairs of vertices. The vertices in our graphs will correspond to variables (whence the common symbol V), and the edges will denote a certain relationship that holds in pairs of variables, the interpretation of which will vary with the application. Two variables connected by an edge are called adjacent.

Each edge in a graph can be either directed (marked by a single arrowhead on the edge), or undirected (unmarked links). In some applications we will also use "bidirected" edges to denote the existence of unobserved common causes (sometimes called *confounders*). These edges will be marked as dotted curved arcs with two arrowheads (see Figure 1.1(a)). If all edges are directed (see Figure 1.1(b)), we then have a *directed* graph. If we strip away all arrowheads from the edges in a graph G, the resultant undirected graph is called the *skeleton* of G. A *path* in a graph is a sequence of edges (e.g., (W, Z), (Z, Y), (Y, X), (X, Z)) in Figure 1.1(a)) such that each edge starts with the vertex ending the preceding edge. In other words, a path is any unbroken, nonintersecting route traced out along the edges in a graph, which may go either along or against the arrows. If every edge in a path is an arrow that points from the first to the second vertex of the pair, we have a *directed path*. In Figure 1.1(a), for example, the path ((W, Z), (Z, Y)) is directed, but the paths ((W, Z), (Z, Y), (Y, X)) and ((W, Z), (Z, X)) are not. If there exists a path between two vertices in a graph, then the two vertices are said to be *connected*; else they are *disconnected*.

Directed graphs may include directed cycles (e.g.,  $X \rightarrow Y$ ,  $Y \rightarrow X$ ), representing mutual causation or feedback processes, but not self-loops (e.g.,  $X \rightarrow X$ ). A graph (like the two in Figure 1.1) that contains no directed cycles is called *acyclic*. A graph that is

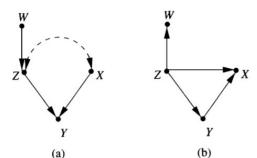


Figure 1.1 (a) A graph containing both directed and bidirected edges. (b) A directed acyclic graph (DAG) with the same skeleton as (a).

both directed and acyclic (Figure 1.1(b)) is called a *directed acyclic graph* (DAG), and such graphs will occupy much of our discussion of causality. We make free use of the terminology of kinship (e.g., *parents*, *children*, *descendants*, *ancestors*, *spouses*) to denote various relationships in a graph. These kinship relations are defined along the full arrows in the graph, including arrows that form directed cycles but ignoring bidirected and undirected edges. In Figure 1.1(a), for example, Y has two parents (X and X), three ancestors (X, X, and X), and no children, while X has no parents (hence, no ancestors), one spouse (X), and one child (X). A *family* in a graph is a set of nodes containing a node and all its parents. For example, X, X, and X, and X, are the families in the graph of Figure 1.1(a).

A node in a directed graph is called a root if it has no parents and a sink if it has no children. Every DAG has at least one root and at least one sink. A connected DAG in which every node has at most one parent is called a tree, and a tree in which every node has at most one child is called a chain. A graph in which every pair of nodes is connected by an edge is called complete. The graph in Figure 1.1(a), for instance, is connected but not complete, because the pairs (W, X) and (W, Y) are not adjacent.

#### 1.2.2 Bayesian Networks

The role of graphs in probabilistic and statistical modeling is threefold:

- 1. to provide convenient means of expressing substantive assumptions;
- 2. to facilitate economical representation of joint probability functions; and
- to facilitate efficient inferences from observations.

We will begin our discussion with item 2.

Consider the task of specifying an arbitrary joint distribution,  $P(x_1,...,x_n)$ , for n dichotomous variables. To store  $P(x_1,...,x_n)$  explicitly would require a table with  $2^n$  entries, an unthinkably large number by any standard. Substantial economy can be achieved when each variable depends on just a small subset of other variables. Such dependence information permits us to decompose large distribution functions into several small distributions – each involving a small subset of variables – and then to piece them together coherently to answer questions of a global nature. Graphs play an essential role in such decomposition, for they provide a vivid representation of the sets of variables that are relevant to each other in any given state of knowledge.

Both directed and undirected graphs have been used by researchers to facilitate such decomposition. Undirected graphs, sometimes called *Markov networks* (Pearl 1988b), are used primarily to represent symmetrical spatial relationships (Isham 1981; Cox and Wermuth 1996; Lauritzen 1996). Directed graphs, especially DAGs, have been used to represent causal or temporal relationships (Lauritzen 1982; Wermuth and Lauritzen 1983; Kiiveri et al. 1984) and came to be known as *Bayesian networks*, a term coined in Pearl (1985) to emphasize three aspects: (1) the subjective nature of the input information; (2) the reliance on Bayes's conditioning as the basis for updating information; and (3) the distinction between causal and evidential modes of reasoning, a distinction that underscores Thomas Bayes's paper of 1763. Hybrid graphs (involving both directed and undirected edges) have also been proposed for statistical modeling (Wermuth and Lauritzen 1990), but in this book our main interest will focus on directed acyclic graphs, with occasional use of directed cyclic graphs to represent feedback cycles.

The basic decomposition scheme offered by directed acyclic graphs can be illustrated as follows. Suppose we have a distribution P defined on n discrete variables, which we may order arbitrarily as  $X_1, X_2, ..., X_n$ . The chain rule of probability calculus (equation (1.12)) always permits us to decompose P as a product of n conditional distributions:

$$P(x_1, \dots, x_n) = \prod_j P(x_j \mid x_1, \dots, x_{j-1}).$$
(1.30)

Now suppose that the conditional probability of some variable  $X_j$  is not sensitive to all the predecessors of  $X_j$  but only to a small subset of those predecessors. In other words, suppose that  $X_j$  is independent of all other predecessors, once we know the value of a select group of predecessors called  $PA_j$ . We can then write

$$P(x_j | x_1, \dots, x_{j-1}) = P(x_j | pa_j)$$
(1.31)

in the product of (1.30), which will considerably simplify the input information required. Instead of specifying the probability of  $X_j$  conditional on all possible realizations of its predecessors  $X_1, \ldots, X_{j-1}$ , we need only concern ourselves with the possible realizations of the set  $PA_j$ . The set  $PA_j$  is called the *Markovian parents* of  $X_j$ , or *parents* for short. The reason for the name becomes clear when we build graphs around this concept.

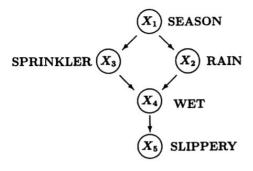
#### **Definition 1.2.1 (Markovian Parents)**

Let  $V = \{X_1, ..., X_n\}$  be an ordered set of variables, and let P(v) be the joint probability distribution on these variables. A set of variables  $PA_j$  is said to be Markovian parents of  $X_j$  if  $PA_j$  is a minimal set of predecessors of  $X_j$  that renders  $X_j$  independent of all its other predecessors. In other words,  $PA_j$  is any subset of  $\{X_1, ..., X_{j-1}\}$  satisfying

$$P(x_j \mid pa_j) = P(x_j \mid x_1, \dots, x_{j-1})$$
(1.32)

and such that no proper subset of PA<sub>i</sub> satisfies (1.32).<sup>5</sup>

<sup>&</sup>lt;sup>5</sup> Lowercase symbols (e.g.,  $x_j$ ,  $pa_j$ ) denote particular realizations of the corresponding variables (e.g.,  $X_j$ ,  $PA_j$ ).



**Figure 1.2** A Bayesian network representing dependencies among five variables.

Definition 1.2.1 assigns to each variable  $X_i$  a select set  $PA_i$  of preceding variables that are sufficient for determining the probability of  $X_i$ ; knowing the values of other preceding variables is redundant once we know the values  $pa_i$  of the parent set  $PA_i$ . This assignment can be represented in the form of a DAG in which variables are represented by nodes and arrows are drawn from each node of the parent set  $PA_i$  toward the child node  $X_i$ . Definition 1.2.1 also suggests a simple recursive method for constructing such a DAG: Starting with the pair  $(X_1, X_2)$ , we draw an arrow from  $X_1$  to  $X_2$  if and only if the two variables are dependent. Continuing to  $X_3$ , we draw no arrow in case  $X_3$  is independent of  $\{X_1, X_2\}$ ; otherwise, we examine whether  $X_2$  screens off  $X_3$  from  $X_1$  or  $X_1$ screens off  $X_3$  from  $X_2$ . In the first case, we draw an arrow from  $X_2$  to  $X_3$ ; in the second, we draw an arrow from  $X_1$  to  $X_3$ . If no screening condition is found, we draw arrows to  $X_3$  from both  $X_1$  and  $X_2$ . In general: at the jth stage of the construction, we select any minimal set of  $X_i$ 's predecessors that screens off  $X_i$  from its other predecessors (as in equation (1.32)), call this set  $PA_i$  and draw an arrow from each member in  $PA_i$  to  $X_i$ . The result is a directed acyclic graph, called a Bayesian network, in which an arrow from  $X_i$  to  $X_i$  assigns  $X_i$ , as a Markovian parent of  $X_i$ , consistent with Definition 1.2.1.

It can be shown (Pearl 1988b) that the set  $PA_j$  is unique whenever the distribution P(v) is strictly positive (i.e., involving no logical or definitional constraints), so that every configuration v of variables, no matter how unlikely, has some finite probability of occurring. Under such conditions, the Bayesian network associated with P(v) is unique, given the ordering of the variables.

Figure 1.2 illustrates a simple yet typical Bayesian network. It describes relationships among the season of the year  $(X_1)$ , whether rain falls  $(X_2)$ , whether the sprinkler is on  $(X_3)$ , whether the pavement would get wet  $(X_4)$ , and whether the pavement would be slippery  $(X_5)$ . All variables in this figure are binary (taking a value of either true or false) except for the root variable  $X_1$ , which can take one of four values: spring, summer, fall, or winter. The network was constructed in accordance with Definition 1.2.1, using causal intuition as a guide. The absence of a direct link between  $X_1$  and  $X_5$ , for example, captures our understanding that the influence of seasonal variations on the slipperiness of the pavement is mediated by other conditions (e.g., the wetness of the pavement). This intuition coincides with the independence condition of  $\{1.32\}$ , since knowing  $X_4$  renders  $X_5$  independent of  $\{X_1, X_2, X_3\}$ .

The construction implied by Definition 1.2.1 defines a Bayesian network as a carrier of conditional independence relationships along the order of construction. Clearly, every distribution satisfying (1.32) must decompose (using the chain rule of (1.30)) into the product

$$P(x_1, ..., x_n) = \prod_{i} P(x_i | pa_i).$$
 (1.33)

For example, the DAG in Figure 1.2 induces the decomposition

$$P(x_1, x_2, x_3, x_4, x_5) = P(x_1)P(x_2 \mid x_1)P(x_3 \mid x_1)P(x_4 \mid x_2, x_3)P(x_5 \mid x_4). \tag{1.34}$$

The product decomposition in (1.33) is no longer order-specific since, given P and G, we can test whether P decomposes into the product given by (1.33) without making any reference to variable ordering. We therefore conclude that a necessary condition for a DAG G to be a Bayesian network of probability distribution P is for P to admit the product decomposition dictated by G, as given in (1.33).

#### **Definition 1.2.2 (Markov Compatibility)**

If a probability function P admits the factorization of (1.33) relative to DAG G, we say that G represents P, that G and P are compatible, or that P is Markov relative to G.

Ascertaining compatibility between DAGs and probabilities is important in statistical modeling primarily because compatibility is a necessary and sufficient condition for a DAG G to explain a body of empirical data represented by P, that is, to describe a stochastic process capable of generating P (e.g., Pearl 1988b, pp. 210–23). If the value of each variable  $X_i$  is chosen at random with some probability  $P_i(x_i \mid pa_i)$ , based solely on the values  $pa_i$  previously chosen for  $PA_i$ , then the overall distribution P of the generated instances  $x_1, x_2, ..., x_n$  will be Markov relative to G. Conversely, if P is Markov relative to G, then there exists a set of probabilities  $P_i(x_i \mid pa_i)$  according to which we can choose the value of each variable  $X_i$  such that the distribution of the generated instances  $x_1, x_2, ..., x_n$  will be equal to P. (In fact, the correct choice of  $P_i(x_i \mid pa_i)$  would be simply  $P(x_i \mid pa_i)$ .)

A convenient way of characterizing the set of distributions compatible with a DAG G is to list the set of (conditional) independencies that each such distribution must satisfy. These independencies can be read off the DAG by using a graphical criterion called d-separation (Pearl 1988b; the d denotes d-rectional), which will play a major role in many discussions in this book.

#### 1.2.3 The d-Separation Criterion

Consider three disjoint sets of variables, X, Y, and Z, which are represented as nodes in a directed acyclic graph G. To test whether X is independent of Y given Z in any distribution compatible with G, we need to test whether the nodes corresponding to variables Z "block" all paths from nodes in X to nodes in Y. By path we mean a sequence of consecutive edges (of any directionality) in the graph, and blocking is to be interpreted as stopping the flow of information (or of dependency) between the variables that are connected by such paths, as defined next.

#### **Definition 1.2.3 (***d***-Separation)**

A path p is said to be d-separated (or blocked) by a set of nodes Z if and only if

<sup>&</sup>lt;sup>6</sup> The latter expression seems to gain strength in recent literature (e.g., Spirtes et al. 1993; Lauritzen 1996). Pearl (1988b, p. 116) used "G is an *I-map* of *P*."

- 1. p contains a chain  $i \to m \to j$  or a fork  $i \leftarrow m \to j$  such that the middle node m is in Z, or
- 2. p contains an inverted fork (or collider)  $i \rightarrow m \leftarrow j$  such that the middle node m is not in Z and such that no descendant of m is in Z.

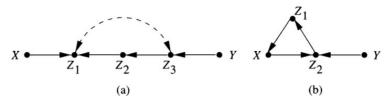
A set Z is said to d-separate X from Y if and only if Z blocks every path from a node in X to a node in Y.

The intuition behind d-separation is simple and can best be recognized if we attribute causal meaning to the arrows in the graph. In causal chains  $i \to m \to j$  and causal forks  $i \leftarrow m \to j$ , the two extreme variables are marginally dependent but become independent of each other (i.e., blocked) once we condition on (i.e., know the value of) the middle variable. Figuratively, conditioning on m appears to "block" the flow of information along the path, since learning about i has no effect on the probability of j, given m. Inverted forks  $i \to m \leftarrow j$ , representing two causes having a common effect, act the opposite way; if the two extreme variables are (marginally) independent, they will become dependent (i.e., connected through unblocked path) once we condition on the middle variable (i.e., the common effect) or any of its descendants. This can be confirmed in the context of Figure 1.2. Once we know the season,  $X_3$  and  $X_2$  are independent (assuming that sprinklers are set in advance, according to the season); whereas finding that the pavement is wet or slippery renders  $X_2$  and  $X_3$  dependent, because refuting one of these explanations increases the probability of the other.

In Figure 1.2,  $X = \{X_2\}$  and  $Y = \{X_3\}$  are d-separated by  $Z = \{X_1\}$ , because both paths connecting  $X_2$  and  $X_3$  are blocked by Z. The path  $X_2 \leftarrow X_1 \rightarrow X_3$  is blocked because it is a fork in which the middle node  $X_1$  is in Z, while the path  $X_2 \rightarrow X_4 \leftarrow X_3$  is blocked because it is an inverted fork in which the middle node  $X_4$  and all its descendants are outside Z. However, X and Y are not d-separated by the set  $Z' = \{X_1, X_5\}$ : the path  $X_2 \rightarrow X_4 \leftarrow X_3$  (an inverted fork) is not blocked by Z', since  $X_5$ , a descendant of the middle node  $X_4$ , is in Z'. 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 arrows converging at  $X_4$ .

At first glance, readers might find it a bit odd that conditioning on a node not lying on a blocked path may unblock the path. However, this corresponds to a general pattern of causal relationships: observations on a common consequence of two independent causes tend to render those causes dependent, because information about one of the causes tends to make the other more or less likely, given that the consequence has occurred. This pattern is known as *selection bias* or *Berkson's paradox* in the statistical literature (Berkson 1946) and as the *explaining away effect* in artificial intelligence (Kim and Pearl 1983). For example, if the admission criteria to a certain graduate school call for either high grades as an undergraduate or special musical talents, then these two attributes will be found to be correlated (negatively) in the student population of that school, even if these attributes are uncorrelated in the population at large. Indeed, students with low grades are likely to be exceptionally gifted in music, which explains their admission to the graduate school.

Figure 1.3 illustrates more elaborate examples of *d*-separation; example (a) contains a bidirected arc  $Z_1 \blacktriangleleft -- \blacktriangleright Z_3$ , and (b) involves a directed cycle  $X \to Z_2 \to Z_1 \to X$ . In



**Figure 1.3** Graphs illustrating d-separation. In (a), X and Y are d-separated given  $Z_2$  and d-connected given  $Z_1$ . In (b), X and Y cannot be d-separated by any set of nodes.

Figure 1.3(a), the two paths between X and Y are blocked when none of  $\{Z_1, Z_2, Z_3\}$  is measured. However, the path  $X \to Z_1 \blacktriangleleft - - \blacktriangleright Z_3 \longleftrightarrow Y$  becomes unblocked when  $Z_1$  is measured. This is so because  $Z_1$  unblocks the "colliders" at both  $Z_1$  and  $Z_3$ ; the first because  $Z_1$  is the collision node of the collider, the second because  $Z_1$  is a descendant of the collision node  $Z_3$  through the path  $Z_1 \leftarrow Z_2 \leftarrow Z_3$ . In Figure 1.3(b), X and Y cannot be d-separated by any set of nodes, including the empty set. If we condition on  $Z_2$ , we block the path  $X \leftarrow Z_1 \leftarrow Z_2 \leftarrow Y$  yet unblock the path  $X \to Z_2 \leftarrow Y$ . If we condition on  $Z_1$ , we again block the path  $X \leftarrow Z_1 \leftarrow Z_2 \leftarrow Y$  and unblock the path  $X \to Z_2 \leftarrow Y$ , because  $Z_1$  is a descendant of the collision node  $Z_2$ .

The connection between *d*-separation and conditional independence is established through the following theorem due to Verma and Pearl (1988; see also Geiger et al. 1990).

#### **Theorem 1.2.4 (Probabilistic Implications of** *d***-Separation)**

If sets X and Y are d-separated by Z in a DAG G, then X is independent of Y conditional on Z in every distribution compatible with G. Conversely, if X and Y are not d-separated by Z in a DAG G, then X and Y are dependent conditional on Z in at least one distribution compatible with G.

The converse part of Theorem 1.2.4 is in fact much stronger – the absence of *d*-separation implies dependence in *almost all* distributions compatible with *G*. The reason is that a precise tuning of parameters is required to generate independency along an unblocked path in the diagram, and such tuning is unlikely to occur in practice (see Spirtes et al. 1993 and Sections 2.4 and 2.9.1).

In order to distinguish between the probabilistic notion of conditional independence  $(X \perp\!\!\!\perp Y \mid Z)_P$  and the graphical notion of *d*-separation, for the latter we will use the notation  $(X \perp\!\!\!\perp Y \mid Z)_G$ . We can thereby express Theorem 1.2.4 more succinctly as follows.

#### Theorem 1.2.5

For any three disjoint subsets of nodes (X,Y,Z) in a DAG G and for all probability functions P, we have:

- (i)  $(X \perp\!\!\!\perp Y \mid Z)_G \Longrightarrow (X \perp\!\!\!\perp Y \mid Z)_P$  whenever G and P are compatible; and
- (ii) if  $(X \perp\!\!\!\perp Y \mid Z)_P$  holds in all distributions compatible with G, it follows that  $(X \perp\!\!\!\perp Y \mid Z)_G$ .

An alternative test for *d*-separation has been devised by Lauritzen et al. (1990), based on the notion of ancestral graphs. To test for  $(X \perp \!\!\! \perp \!\!\! \perp \!\!\! \perp \!\!\! \mid Z)_G$ , delete from *G* all nodes except those in  $\{X,Y,Z\}$  and their ancestors, connect by an edge every pair of nodes that share

a common child, and remove all arrows from the arcs. Then  $(X \perp\!\!\!\perp Y \mid Z)_G$  holds if and only if Z intercepts all paths between X and Y in the resulting undirected graph.

Note that the ordering with which the graph was constructed does not enter into the d-separation criterion; it is only the topology of the resulting graph that determines the set of independencies that the probability P must satisfy. Indeed, the following theorem can be proven (Pearl 1988b, p. 120).

#### Theorem 1.2.6 (Ordered Markov Condition)

A necessary and sufficient condition for a probability distribution P to be Markov relative a DAG G is that, conditional on its parents in G, each variable be independent of all its predecessors in some ordering of the variables that agrees with the arrows of G.

A consequence of this theorem is an order-independent criterion for determining whether a given probability P is Markov relative to a given DAG G.

#### **Theorem 1.2.7 (Parental Markov Condition)**

A necessary and sufficient condition for a probability distribution P to be Markov relative a DAG G is that every variable be independent of all its nondescendants (in G), conditional on its parents. (We exclude  $X_i$  when speaking of its "nondescendants.")

This condition, which Kiiveri et al. (1984) and Lauritzen (1996) called the "local" Markov condition, is sometimes taken as the definition of Bayesian networks (Howard and Matheson 1981). In practice, however, the ordered Markov condition is easier to use.

Another important property that follows from d-separation is a criterion for determining whether two given DAGs are observationally equivalent – that is, whether every probability distribution that is compatible with one of the DAGs is also compatible with the other.

#### **Theorem 1.2.8 (Observational Equivalence)**

Two DAGs are observationally equivalent if and only if they have the same skeletons and the same sets of v-structures, that is, two converging arrows whose tails are not connected by an arrow (Verma and Pearl 1990).<sup>7</sup>

Observational equivalence places a limit on our ability to infer directionality from probabilities alone. Two networks that are observationally equivalent cannot be distinguished without resorting to manipulative experimentation or temporal information. For example, reversing the direction of the arrow between  $X_1$  and  $X_2$  in Figure 1.2 would neither introduce nor destroy a v-structure. Therefore, this reversal yields an observationally equivalent network, and the directionality of the link  $X_1 \rightarrow X_2$  cannot be determined from probabilistic information. The arrows  $X_2 \rightarrow X_4$  and  $X_4 \rightarrow X_5$ , however, are of different nature; there is no way of reversing their directionality without creating a new v-structure. Thus, we see that some probability functions P (such as the one responsible for the construction of the Bayesian network in Figure 1.2), when unaccompanied

An identical criterion was independently derived by Frydenberg (1990) in the context of chain graphs, where strict positivity is assumed.

by temporal information, can constrain the directionality of some arrows in the graph. The precise meaning of such directionality constraints – and the possibility of using these constraints for inferring causal relationships from data – will be formalized in Chapter 2.

## 1.2.4 Inference with Bayesian Networks

Bayesian networks were developed in the early 1980s to facilitate the tasks of prediction and "abduction" in artificial intelligence (AI) systems. In these tasks, it is necessary to find a coherent interpretation of incoming observations that is consistent with both the observations and the prior information at hand. Mathematically, the task boils down to the computation of  $P(y \mid x)$ , where X is a set of observations and Y is a set of variables that are deemed important for prediction or diagnosis.

Given a joint distribution P, the computation of  $P(y \mid x)$  is conceptually trivial and invokes straightforward application of Bayes's rule to yield

$$P(y \mid x) = \frac{\sum_{s} P(y, x, s)}{\sum_{y, s} P(y, x, s)},$$
(1.35)

where S stands for the set of all variables excluding X and Y. Because every Bayesian network defines a joint probability P (given by the product in (1.33)), it is clear that  $P(y \mid x)$  can be computed from a DAG G and the conditional probabilities  $P(x_i \mid pa_i)$  defined on the families of G.

The challenge, however, lies in performing these computations efficiently and within the representation level provided by the network topology. The latter is important in systems that generate explanations for their reasoning processes. Although such inference techniques are not essential to our discussion of causality, we will nevertheless survey them briefly, for they demonstrate (i) the effectiveness of organizing probabilistic knowledge in the form of graphs and (ii) the feasibility of performing coherent probabilistic calculations (and approximations thereof) on such organization. Details can be found in the references cited.

The first algorithms proposed for probabilistic calculations in Bayesian networks used message-passing architecture and were limited to trees (Pearl 1982; Kim and Pearl 1983). With this technique, each variable is assigned a simple processor and permitted to pass messages asynchronously to its neighbors until equilibrium is achieved (in a finite number of steps). Methods have since been developed that extend this tree propagation (and some of its synchronous variants) to general networks. Among the most popular are Lauritzen and Spiegelhalter's (1988) method of join-tree propagation and the method of cut-set conditioning (Pearl 1988b, pp. 204–10; Jensen 1996). In the join-tree method, we decompose the network into clusters (e.g., cliques) that form tree structures and then treat the set variables in each cluster as a compound variable that is capable of passing messages to its neighbors (which are also compound variables). For example, the network of Figure 1.2 can be structured as a Markov-compatible chain of three clusters:

$${X_1, X_2, X_3} \rightarrow {X_2, X_3, X_4} \rightarrow {X_4, X_5}.$$

In the cut-set conditioning method, a set of variables is instantiated (given specific values) such that the remaining network forms a tree. The propagation is then performed on that tree, and a new instantiation chosen, until all instantiations have been exhausted; the results are then averaged. In Figure 1.2, for example, if we instantiate  $X_1$  to any specific value (say,  $X_1$  = summer), then we break the pathway between  $X_2$  and  $X_3$  and the remaining network becomes tree-structured. The main advantage of the cut-set conditioning method is that its storage-space requirement is minimal (linear in the size of the network), whereas that of the join-tree method might be exponential. Hybrid combinations of these two basic algorithms have also been proposed (Shachter et al. 1994; Dechter 1996) to allow flexible trade-off of storage versus time (Darwiche 2009).

Whereas inference in general networks is "NP-hard" (Cooper 1990), the computational complexity for each of the methods cited here can be estimated prior to actual processing. When the estimates exceed reasonable bounds, an approximation method such as stochastic simulation (Pearl 1988b, pp. 210–23) can be used instead. This method exploits the topology of the network to perform Gibbs sampling on local subsets of variables, sequentially as well as concurrently.

Additional properties of DAGs and their applications to evidential reasoning in expert systems are discussed in Pearl (1988b), Lauritzen and Spiegelhalter (1988), Pearl (1993a), Spiegelhalter et al. (1993), Heckerman et al. (1995), and Darwiche (2009).

#### 1.3 CAUSAL BAYESIAN NETWORKS

The interpretation of direct acyclic graphs as carriers of independence assumptions does not necessarily imply causation; in fact, it will be valid for any set of recursive independencies along any ordering of the variables, not necessarily causal or chronological. However, the ubiquity of DAG models in statistical and AI applications stems (often unwittingly) primarily from their causal interpretation – that is, as a system of processes, one per family, that could account for the generation of the observed data. It is this causal interpretation that explains why DAG models are rarely used in any variable ordering other than those which respect the direction of time and causation.

The advantages of building DAG models around causal rather than associational information are several. First, the judgments required in the construction of the model are more meaningful, more accessible and hence more reliable. The reader may appreciate this point by attempting to construct a DAG representation for the associations in Figure 1.2 along the ordering  $(X_5, X_1, X_3, X_2, X_4)$ . Such exercises illustrate not only that some independencies are more vividly accessible to the mind than others but also that conditional independence judgments are accessible (hence reliable) only when they are anchored onto more fundamental building blocks of our knowledge, such as causal relationships. In the example of Figure 1.2, our willingness to assert that  $X_5$  is independent of  $X_2$  and  $X_3$  once we know  $X_4$  (i.e., whether the pavement is wet) is defensible because we can easily translate the assertion into one involving causal relationships: that the *influence* of rain and sprinkler on slipperiness is *mediated* by the wetness of the pavement. Dependencies that are not supported by causal links are considered odd or spurious and are even branded "paradoxical" (see the discussion of Berkson's paradox, Section 1.2.3).

We will have several opportunities throughout this book to demonstrate the primacy of causal over associational knowledge. In extreme cases, we will see that people tend to ignore probabilistic information altogether and attend to causal information instead (see Section 6.1.4). This puts into question the ruling paradigm of graphical models in statistics (Wermuth and Lauritzen 1990; Cox and Wermuth 1996), according to which conditional independence assumptions are the primary vehicle for expressing substantive knowledge. It seems that if conditional independence judgments are by-products of stored causal relationships, then tapping and representing those relationships directly would be a more natural and more reliable way of expressing what we know or believe about the world. This is indeed the philosophy behind causal Bayesian networks.

The second advantage of building Bayesian networks on causal relationships – one that is basic to the understanding of causal organizations – is the ability to represent and respond to external or spontaneous *changes*. Any local reconfiguration of the mechanisms in the environment can be translated, with only minor modification, into an isomorphic reconfiguration of the network topology. For example, to represent a disabled sprinkler in the story of Figure 1.2, we simply delete from the network all links incident to the node Sprinkler. To represent the policy of turning the sprinkler off if it rains, we simply add a link between Rain and Sprinkler and revise  $P(x_3 \mid x_1, x_2)$ . Such changes would require much greater remodeling efforts if the network were not constructed along the causal direction but instead along (say) the order  $(X_5, X_1, X_3, X_2, X_4)$ . This remodeling flexibility may well be cited as the ingredient that marks the division between deliberative and reactive agents and that enables the former to manage novel situations instantaneously, without requiring training or adaptation.

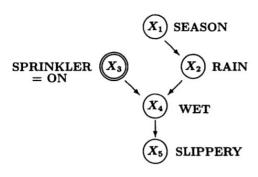
## 1.3.1 Causal Networks as Oracles for Interventions

The source of this flexibility rests on the assumption that each parent—child relationship in the network represents a stable and autonomous physical mechanism – in other words, that it is conceivable to change one such relationship *without* changing the others. Organizing one's knowledge in such *modular* configurations permits one to predict the effect of external interventions with a minimum of extra information. Indeed, causal models (assuming they are valid) are much more informative than probability models. 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 – such as those encountered in policy analysis, treatment management, or planning everyday activity. Such changes cannot be deduced from a joint distribution, even if fully specified.

The connection between modularity and interventions is as follows. Instead of specifying a new probability function for each of the many possible interventions, we specify

<sup>8</sup> The Tversky and Kahneman (1980) experiments with causal biases in probability judgment constitute another body of evidence supporting this observation. For example, most people believe that it is more likely for a girl to have blue eyes, given that her mother has blue eyes, than the other way around; the two probabilities are in fact equal.

<sup>&</sup>lt;sup>9</sup> The author was as guilty of advocating the centrality of conditional independence as were his colleagues in statistics; see Pearl (1988b, p. 79).



**Figure 1.4** Network representation of the action "turning the sprinkler On."

merely the immediate change implied by the intervention and, by virtue of autonomy, we assume that the change is local, and does not spread over to mechanisms other than those specified. Once we know the identity of the mechanism altered by an intervention and the nature of the alteration, the overall effect of an intervention can be predicted by modifying the corresponding factors in (1.33) and using the modified product to compute a new probability function. For example, to represent the action "turning the sprinkler On" in the network of Figure 1.2, we delete the link  $X_1 \rightarrow X_3$  and assign  $X_3$  the value On. The graph resulting from this operation is shown in Figure 1.4, and the resulting joint distribution on the remaining variables will be

$$P_{X_3=\text{On}}(x_1, x_2, x_4, x_5) = P(x_1) P(x_2 \mid x_1) P(x_4 \mid x_2, X_3 = \text{On}) P(x_5 \mid x_4),$$
 (1.36)

in which all the factors on the right-hand side (r.h.s.), by virtue of autonomy, are the same as in (1.34).

The deletion of the factor  $P(x_3 \mid x_1)$  represents the understanding that, whatever relationship existed between seasons and sprinklers prior to the action, that relationship is no longer in effect while we perform the action. Once we physically turn the sprinkler on and keep it on, a new mechanism (in which the season has no say) determines the state of the sprinkler.

Note the difference between the action  $do(X_3 = \text{On})$  and the observation  $X_3 = \text{On}$ . The effect of the latter is obtained by ordinary Bayesian conditioning, that is,  $P(x_1, x_2, x_4, x_5 | X_3 = \text{On})$ , while that of the former by conditioning a mutilated graph, with the link  $X_1 \rightarrow X_3$  removed. This indeed mirrors the difference between seeing and doing: after *observing* that the sprinkler is on, we wish to infer that the season is dry, that it probably did not rain, and so on; no such inferences should be drawn in evaluating the effects of a contemplated *action* "turning the sprinkler On."

The ability of causal networks to predict the effects of actions of course requires a stronger set of assumptions in the construction of those networks, assumptions that rest on causal (not merely associational) knowledge and that ensure the system would respond to interventions in accordance with the principle of autonomy. These assumptions are encapsulated in the following definition of causal Bayesian networks.

#### **Definition 1.3.1 (Causal Bayesian Network)**

Let P(v) be a probability distribution on a set V of variables, and let  $P_x(v)$  denote the distribution resulting from the intervention do(X = x) that sets a subset X of variables

to constants x.<sup>10</sup> Denote by  $P_*$  the set of all interventional distributions  $P_X(v)$ ,  $X \subseteq V$ , including P(v), which represents no intervention (i.e.,  $X = \phi$ ). A DAG G is said to be a causal Bayesian network compatible with  $P_*$  if and only if the following three conditions hold for every  $P_X \in P_*$ :

- (i)  $P_r(v)$  is Markov relative to G;
- (ii)  $P_x(v_i) = 1$  for all  $V_i \in X$  whenever  $v_i$  is consistent with X = x;
- (iii)  $P_x(v_i|pa_i) = P(v_i|pa_i)$  for all  $V_i \notin X$  whenever  $pa_i$  is consistent with X = x, i.e., each  $P(v_i|pa_i)$  remains invariant to interventions not involving  $V_i$ .

Definition 1.3.1 imposes constraints on the interventional space  $P_*$  that permit us to encode this vast space economically, in the form of a single Bayesian network G. These constraints enable us to compute the distribution  $P_x(v)$  resulting from any intervention do(X = x) as a truncated factorization

$$P_{x}(v) = \prod_{\{i \mid V_{i} \notin X\}} P(v_{i} \mid pa_{i}) \qquad \text{for all } v \text{ consistent with } x,$$

$$(1.37)$$

which follows from (and implies) conditions (i)–(iii), thus justifying the family deletion procedure on G, as in (1.36). It is not hard to show that, whenever G is a causal Bayes network with respect to  $P_*$ , the following two properties must hold.

## **Property 1**

For all i,

$$P(v_i \mid pa_i) = P_{pa_i}(v_i). \tag{1.38}$$

#### Property 2

For all i and for every subset S of variables disjoint of  $\{V_i, PA_i\}$ , we have

$$P_{pa_{i},s}(v_{i}) = P_{pa_{i}}(v_{i}). {(1.39)}$$

Property 1 renders every parent set  $PA_i$  exogenous relative to its child  $V_i$ , ensuring that the conditional probability  $P(v_i | pa_i)$  coincides with the effect (on  $V_i$ ) of setting  $PA_i$  to  $pa_i$  by external control. Property 2 expresses the notion of invariance; once we control its direct causes  $PA_i$ , no other interventions will affect the probability of  $V_i$ .

## 1.3.2 Causal Relationships and Their Stability

This mechanism-based conception of interventions provides a semantical basis for notions such as "causal effects" or "causal influence," to be defined formally and analyzed in Chapters 3 and 4. For example, to test whether a variable  $X_i$  has a causal influence on another variable  $X_j$ , we compute (using the truncated factorization formula of (1.37)) the (marginal) distribution of  $X_j$  under the actions  $do(X_i = x_i)$  – namely,  $P_{X_i}(x_j)$  for all

<sup>&</sup>lt;sup>10</sup> The notation  $P_x(v)$  will be replaced in subsequent chapters with  $P(v \mid do(x))$  and  $P(v \mid \hat{x})$  to facilitate algebraic manipulations.

values  $x_i$  of  $X_i$  – and test whether that distribution is sensitive to  $x_i$ . It is easy to see from our previous examples that only variables that are descendants of  $X_i$  in the causal network can be influenced by  $X_i$ ; deleting the factor  $P(x_i|pa_i)$  from the joint distribution turns  $X_i$  into a root node in the mutilated graph, and root variables (as the d-separation criterion dictates) are independent of all other variables except their descendants.

This understanding of causal influence permits us to see precisely why, and in what way, causal relationships are more "stable" than probabilistic relationships. We expect such difference in stability because causal relationships are *ontological*, describing objective physical constraints in our world, whereas probabilistic relationships are *epistemic*, reflecting what we know or believe about the world. Therefore, causal relationships should remain unaltered as long as no change has taken place in the environment, even when our knowledge about the environment undergoes changes. To demonstrate, consider the causal relationship  $S_1$ , "Turning the sprinkler on would not affect the rain," and compare it to its probabilistic counterpart  $S_2$ , "The state of the sprinkler is independent of (or unassociated with) the state of the rain." Figure 1.2 illustrates two obvious ways in which  $S_2$  will change while  $S_1$  remains intact. First,  $S_2$  changes from false to true when we learn what season it is  $(X_1)$ . Second, given that we know the season,  $S_2$  changes from true to false once we observe that the pavement is wet  $(X_4 = \text{true})$ . On the other hand,  $S_1$  remains true regardless of what we learn or know about the season or about the pavement.

The example reveals a stronger sense in which causal relationships are more stable than the corresponding probabilistic relationships, a sense that goes beyond their basic ontological–epistemological difference. The relationship  $S_1$  will remain invariant to changes in the mechanism that regulates how seasons affect sprinklers. In fact, it remains invariant to changes in *all* mechanisms shown in this causal graph. We thus see that causal relationships exhibit greater robustness to ontological changes as well; they are sensitive to a smaller set of mechanisms. More specifically, and in marked contrast to probabilistic relationships, causal relationships remain invariant to changes in the mechanism that governs the causal variables ( $X_3$  in our example).

In view of this stability, it is no wonder that people prefer to encode knowledge in causal rather than probabilistic structures. Probabilistic relationships, such as marginal and conditional independencies, may be helpful in hypothesizing initial causal structures from uncontrolled observations. However, once knowledge is cast in causal structure, those probabilistic relationships tend to be forgotten; whatever judgments people express about conditional independencies in a given domain are derived from the causal structure acquired. This explains why people feel confident asserting certain conditional independencies (e.g., that the price of beans in China is independent of the traffic in Los Angeles) having no idea whatsoever about the numerical probabilities involved (e.g., whether the price of beans will exceed \$10 per bushel).

The element of stability (of mechanisms) is also at the heart of the so-called explanatory accounts of causality, according to which causal models need not encode behavior under intervention but instead aim primarily to provide an "explanation" or "understanding" of how data are generated.<sup>11</sup> Regardless of what use is eventually made

Elements of this explanatory account can be found in the writings of Dempster (1990), Cox (1992), and Shafer (1996); see also King et al. (1994, p. 75).

of our "understanding" of things, we surely would prefer an understanding in terms of durable relationships, transportable across situations, over those based on transitory relationships. The sense of "comprehensibility" that accompanies an adequate explanation is a natural by-product of the transportability of (and hence of our familiarity with) the causal relationships used in the explanation. It is for reasons of stability that we regard the falling barometer as predicting but not explaining the rain; those predictions are not transportable to situations where the pressure surrounding the barometer is controlled by artificial means. True understanding enables predictions in such novel situations, where some mechanisms change and others are added. It thus seems reasonable to suggest that, in the final analysis, the explanatory account of causation is merely a variant of the manipulative account, albeit one where interventions are dormant. Accordingly, we may as well view our unsatiated quest for understanding "how data is generated" or "how things work" as a quest for acquiring the ability to make predictions under a wider range of circumstances, including circumstances in which things are taken apart, reconfigured, or undergo spontaneous change.

## 1.4 FUNCTIONAL CAUSAL MODELS

The way we have introduced the causal interpretation of Bayesian networks represents a fundamental departure from the way causal models (and causal graphs) were first introduced into genetics (Wright 1921), econometrics (Haavelmo 1943), and the social sciences (Duncan 1975), as well as from the way causal models are used routinely in physics and engineering. In those models, causal relationships are expressed in the form of deterministic, *functional* equations, and probabilities are introduced through the assumption that certain variables in the equations are unobserved. This reflects Laplace's (1814) conception of natural phenomena, according to which nature's laws are deterministic and randomness surfaces owing merely to our ignorance of the underlying boundary conditions. In contrast, all relationships in the definition of causal Bayesian networks were assumed to be inherently stochastic and thus appeal to the modern (i.e., quantum mechanical) conception of physics, according to which all nature's laws are inherently probabilistic and determinism is but a convenient approximation.

In this book, we shall express preference toward Laplace's quasi-deterministic conception of causality and will use it, often contrasted with the stochastic conception, to define and analyze most of the causal entities that we study. This preference is based on three considerations. First, the Laplacian conception is more general. Every stochastic model can be emulated by many functional relationships (with stochastic inputs), but not the other way around; functional relationships can only be approximated, as a limiting case, using stochastic models. Second, the Laplacian conception is more in tune with human intuition. The few esoteric quantum mechanical experiments that conflict with the predictions of the Laplacian conception evoke surprise and disbelief, and they demand that physicists give up deeply entrenched intuitions about locality and causality (Maudlin 1994). Our objective is to preserve, explicate, and satisfy – not destroy – those intuitions. <sup>12</sup>

The often heard argument that human intuitions belong in psychology and not in science or philosophy is inapplicable when it comes to causal intuition – the original authors of causal thoughts

Finally, certain concepts that are ubiquitous in human discourse can be defined only in the Laplacian framework. We shall see, for example, that such simple concepts as "the probability that event B occured because of event A" and "the probability that event B would have been different if it were not for event A" cannot be defined in terms of purely stochastic models. These so-called counterfactual concepts will require a synthesis of the deterministic and probabilistic components embodied in the Laplacian model.

#### 1.4.1 Structural Equations

In its general form, a functional causal model consists of a set of equations of the form

$$x_i = f_i(pa_i, u_i), \quad i = 1, \dots, n,$$
 (1.40)

where  $pa_i$  (connoting *parents*) stands for the set of variables that directly determine the value of  $X_i$  and where the  $U_i$  represent errors (or "disturbances") due to omitted factors. Equation (1.40) is a nonlinear, nonparametric generalization of the linear structural equation models (SEMs)

$$x_i = \sum_{k \neq 1} \alpha_{ik} x_k + u_i, \quad i = 1, \dots, n,$$
 (1.41)

which have become a standard tool in economics and social science (see Chapter 5 for a detailed exposition of this enterprise). In linear models,  $pa_i$  corresponds to those variables on the r.h.s. of (1.41) that have nonzero coefficients.

The interpretation of the functional relationship in (1.40) is the standard interpretation that functions carry in physics and the natural sciences; it is a recipe, a strategy, or a law specifying what value nature would assign to  $X_i$  in response to every possible value combination that  $(PA_i, U_i)$  might take on. A set of equations in the form of (1.40) and in which each equation represents an autonomous mechanism is called a *structural model*; if each variable has a distinct equation in which it appears on the left-hand side (called the *dependent* variable), then the model is called a *structural causal model* or a *causal model* for short. Mathematically, the distinction between structural and algebraic equations is that any subset of structural equations is, in itself, a valid structural model – one that represents conditions under some set of interventions.

To illustrate, Figure 1.5 depicts a canonical econometric model relating price and demand through the equations

$$q = b_1 p + d_1 i + u_1, (1.42)$$

$$p = b_2 q + d_2 w + u_2, (1.43)$$

where Q is the quantity of household demand for a product A, P is the unit price of product A, I is household income, W is the wage rate for producing product A, and  $U_1$  and

cannot be ignored when the meaning of the concept is in question. Indeed, compliance with human intuition has been the ultimate criterion of adequacy in every philosophical study of causation, and the proper incorporation of background information into statistical studies likewise relies on accurate interpretation of causal judgment.

Formal treatment of causal models, structural equations, and error terms are given in Chapter 5 (Section 5.4.1) and Chapter 7 (Sections 7.1 and 7.2.5).

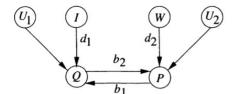


Figure 1.5 Causal diagram illustrating the relationship between price (P), demand (Q), income (I), and wages (W).

 $U_2$  represent error terms – unmodeled factors that affect quantity and price, respectively (Goldberger 1992). The graph associated with this model is cyclic, and the vertices associated with the variables  $U_1$ ,  $U_2$ , I, and W are root nodes, conveying the assumption of mutual independence. The idea of autonomy (Aldrich 1989), in this context, means that the two equations represent two loosely coupled segments of the economy, consumers and producers. Equation (1.42) describes how consumers decide what quantity Q to buy, and (1.43) describes how manufacturers decide what price P to charge. Like all feedback systems, this too represents implicit dynamics; today's prices are determined on the basis of yesterday's demand, and these prices will determine the demand in the next period of transactions. The solution to such equations represents a long-term equilibrium under the assumption that the background quantities,  $U_1$  and  $U_2$ , remain constant.

The two equations are considered to be "autonomous" relative to the dynamics of changes in the sense that external changes affecting one equation do not imply changes to the others. For example, if government decides on price control and sets the price P at  $p_0$ , then (1.43) will be modified to read  $p = p_0$  but the relationships in (1.42) will remain intact, yielding  $q = b_1 p_0 + d_1 i + u_1$ . We thus see that  $b_1$ , the "demand elasticity," should be interpreted as the rate of change of Q per unit controlled change in P. This is different, of course, from the rate of change of Q per unit observed change in P (under uncontrolled conditions), which, besides  $b_1$ , is also affected by the parameters of (1.43) (see Section 7.2.1, equation (7.14)). The difference between controlled and observed changes is essential for the correct interpretation of structural equation models in social science and economics, and it will be discussed at length in Chapter 5. If we have reasons to believe that consumer behavior will also change under a price control policy, then this modified behavior would need to be modeled explicitly – for example, by treating the coefficients  $b_1$  and  $d_1$  as dependent variables in auxiliary equations involving P. 14 Section 7.2.1 will present an analysis of policy-related problems using this model.

To illustrate the workings of nonlinear functional models, consider again the causal relationships depicted in Figure 1.2. The causal model associated with these relationships will consist of five functions, each representing an autonomous mechanism governing one variable:

$$x_1 = u_1,$$
  
 $x_2 = f_2(x_1, u_2),$ 

<sup>&</sup>lt;sup>14</sup> Indeed, consumers normally react to price fixing by hoarding goods in anticipation of shortages (Lucas 1976). Such phenomena are not foreign to structural models, though; they simply call for more elaborate equations to capture consumers' expectations.

of the error distributions  $P(u_i)$ . Thus, we need not specify in advance the functional form of  $\{f_i\}$  or the distributions  $P(u_i)$ ; once we measure (or estimate)  $P(x_i | pa_i)$ , all probabilistic properties of a Markovian causal model are determined, regardless of the mechanism that actually generates those conditional probabilities. Druzdzel and Simon (1993) showed that, for every Bayesian network G characterized by a distribution P (as in (1.33)), there exists a functional model (as in (1.40)) that generates a distribution identical to P. It follows that in all probabilistic applications of Bayesian networks – including statistical estimation, prediction, and diagnosis – we can use an equivalent functional model as specified in (1.40), and we can regard functional models as just another way of encoding joint distribution functions.

Nonetheless, the causal-functional specification has several advantages over the probabilistic specification, even in purely predictive (i.e., nonmanipulative) tasks, First and foremost, all the conditional independencies that are displayed by the causal diagram G are guaranteed to be stable – that is, invariant to parametric changes in the mechanisms represented by the functions  $f_i$  and the distributions  $P(u_i)$ . This means that agents who choose to organize knowledge using Markovian causal models can make reliable assertions about conditional independence relations without assessing numerical probabilities – a common ability among humanoids <sup>19</sup> and a useful feature for inference. Second, the functional specification is often more meaningful and natural, and it yields a small number of parameters. Typical examples are the linear structural equations used in social science and economics (see Chapter 5) and the "noisy OR gate" that has become quite popular in modeling the effect of multiple dichotomous causes (Pearl 1988b, p. 184). Third (and perhaps hardest for an empiricist to accept), judgmental assumptions of conditional independence among observable quantities are simplified and made more reliable in functional models, because such assumptions are cast directly as judgments about the presence or absence of *unobserved* common causes (e.g., why is the price of beans in China judged to be independent of the traffic in Los Angeles?). In the construction of Bayesian networks, for example, instead of judging whether each variable is independent of all its nondescendants (given its parents), we need to judge whether the parent set contains all relevant immediate causes – in particular, whether no factor omitted from the parent set is a cause of another observed variable. Such judgments are more natural because they are discernible directly from a qualitative causal structure, the very structure that our mind has selected for storing stable aspects of experience.

Finally, there is an additional advantage to basing prediction models on causal mechanisms that stems from considerations of stability (Section 1.3.2). When some conditions in the environment undergo change, it is usually only a few causal mechanisms that are affected by the change; the rest remain unaltered. It is simpler then to reassess (judgmentally) or reestimate (statistically) the model parameters knowing that

<sup>&</sup>lt;sup>18</sup> In Chapter 9 we will show that, except in some pathological cases, there actually exist an infinite number of functional models with this property.

Statisticians who are reluctant to discuss causality yet have no hesitation expressing background information in the form of conditional independence statements would probably be shocked to realize that such statements acquire their validity from none other than the *causal* Markov condition (Theorem 1.4.1). See note 9.

the corresponding symbolic change is also local, involving just a few parameters, than to reestimate the entire model from scratch.<sup>20</sup>

#### 1.4.3 Interventions and Causal Effects in Functional Models

The functional characterization  $x_i = f_i(pa_i, u_i)$ , like its stochastic counterpart, provides a convenient language for specifying how the resulting distribution would change in response to external interventions. This is accomplished by encoding each intervention as an alteration on a select set of functions instead of a select set of conditional probabilities. The overall effect of the intervention can then be predicted by modifying the corresponding equations in the model and using the modified model to compute a new probability function. Thus, all features of causal Bayesian networks (Section 1.3) can be emulated in Markovian functional models.

For example, to represent the action "turning the sprinkler On" in the model of (1.44), we delete the equation  $x_3 = f_3(x_1, u_3)$  and replace it with  $x_3 = On$ . The modified model will contain all the information needed for computing the effect of the action on other variables. For example, the probability function induced by the modified model will be equal to that given by (1.36), and the modified diagram will coincide with that of Figure 1.4.

More generally, when an intervention forces a subset X of variables to attain fixed values x, then a subset of equations is to be pruned from the model in (1.40), one for each member of X, thus defining a new distribution over the remaining variables that characterizes the effect of the intervention and coincides with the truncated factorization obtained by pruning families from a causal Bayesian network (equation (1.37)).

The functional model's representation of interventions offers greater flexibility and generality than that of a stochastic model. First, the analysis of interventions can be extended to cyclic models, like the one in Figure 1.5, so as to answer policy-related questions  $^{22}$  (e.g.: What would the demand quantity be if we control the price at  $p_0$ ?). Second, interventions involving the modification of equational parameters (like  $b_1$  and  $d_1$  in (1.42)) are more readily comprehended than those described as modifiers of conditional probabilities, perhaps because stable physical mechanisms are normally associated with equations and not with conditional probabilities. Conditional probabilities are perceived to be derivable from, not generators of, joint distributions. Third, the analysis of causal effects in non-Markovian models will be greatly simplified using functional models. The reason is: there are infinitely many conditional probabilities  $P(x_i \mid pa_i)$  but only a finite number of functions  $x_i = f_i(pa_i, u_i)$  among discrete variables  $X_i$  and  $PA_i$ . This fact will enable us in Chapter 8 (Section 8.2.2) to use linear-programming techniques to obtain sharp bounds on causal effects in studies involving noncompliance.

To the best of my knowledge, this aspect of causal models has not been studied formally; it is suggested here as a research topic for students of adaptive systems.

An explicit translation of interventions to "wiping out" equations from the model was first proposed by Strotz and Wold (1960) and later used in Fisher (1970) and Sobel (1990). More elaborate types of interventions, involving conditional actions and stochastic strategies, will be formulated in Chapter 4.

<sup>22</sup> Such questions, especially those involving the control of endogenous variables, are conspicuously absent from econometric textbooks (see Chapter 5).

Finally, functional models permit the analysis of context-specific actions and policies. The notion of causal effect as defined so far is of only minor use in practical policy making. The reason is that causal effects tell us the general tendency of an action to bring about a response (as with the tendency of a drug to enhance recovery in the overall population) but are not specific to actions in a given situation characterized by a set of particular observations that may themselves be affected by the action. A physician is usually concerned with the effect of a treatment on a patient who has already been examined and found to have certain symptoms. Some of those symptoms will themselves be affected by the treatment. Likewise, an economist is concerned with the effect of taxation in a given economic context characterized by various economical indicators, which (again) will be affected by taxation if applied. Such context-specific causal effects cannot be computed by simulating an intervention in a static Bayesian network, because the context itself varies with the intervention and so the conditional probabilities  $P(x_i \mid pa_i)$ are altered in the process. However, the functional relationships  $x_i = f_i(pa_i, u_i)$  remain invariant, which enables us to compute context-specific causal effects as outlined in the next section (see Sections 7.2.1, 8.3, and 9.3.4 for full details).

#### 1.4.4 Counterfactuals in Functional Models

We now turn to the most distinctive characteristic of functional models – the analysis of *counterfactuals*. Certain counterfactual sentences, as we remarked before, cannot be defined in the framework of stochastic causal networks. To see the difficulties, let us consider the simplest possible causal Bayesian network consisting of a pair of independent (hence unconnected) binary variables X and Y. Such a network ensues, for example, in a controlled (i.e., randomized) clinical trial when we find that a treatment X has no effect on the distribution of subjects' response Y, which may stand for either recovery (Y = 0) or death (Y = 1). Assume that a given subject, Joe, has taken the treatment and died; we ask whether Joe's death occurred *because of* the treatment, *despite* the treatment, or *regardless of* the treatment. In other words, we ask for the probability Q that Joe would have died had he not been treated.

To highlight the difficulty in answering such counterfactual questions, let us take an extreme case where 50% of the patients recover and 50% die in both the treatment and the control groups; assume further that the sample size approaches infinity, thus yielding

$$P(y \mid x) = 1/2 \text{ for all } x \text{ and } y. \tag{1.46}$$

Readers versed in statistical testing will recognize immediately the impossibility of answering the counterfactual question from the available data, noting that Joe, who took the treatment and died, was never tested under the no-treatment condition. Moreover, the difficulty does not stem from addressing the question to a particular individual, Joe, for whom we have only one data point. Rephrasing the question in terms of population frequencies – asking what percentage Q of subjects who died under treatment would have recovered had they not taken the treatment – will encounter the same difficulties because none of those subjects was tested under the no-treatment condition. Such difficulties have prompted some statisticians to dismiss counterfactual questions as metaphysical and to

advocate the restriction of statistical analysis to only those questions that can be answered by direct tests (Dawid 2000).

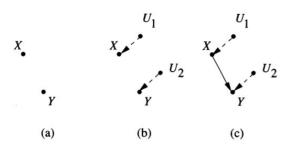
However, that our scientific, legal, and ordinary languages are loaded with counterfactual utterances indicates clearly that counterfactuals are far from being metaphysical; they must have definite testable implications and must carry valuable substantive information. The analysis of counterfactuals therefore represents an opportunity to anyone who shares the aims of this book: integrating substantive knowledge with statistical data so as to refine the former and interpret the latter. Within this framework, the counterfactual issue demands answers to tough, yet manageable technical questions: What is the empirical content of counterfactual queries? What knowledge is required to answer those queries? How can this knowledge be represented mathematically? Given such representation, what mathematical machinery is needed for deriving the answers?

Chapter 7 (Section 7.2.2) presents an empirical explication of counterfactuals as claims about the temporal persistence of certain mechanisms. In our example, the response to treatment of each (surviving) patient is assumed to be persistent. If the outcome Y were a reversible condition, rather than death, then the counterfactual claim would translate directly into predictions about response to future treatments. But even in the case of death, the counterfactual quantity Q implies not merely a speculation about the hypothetical behavior of subjects who died but also a testable claim about surviving untreated subjects under subsequent treatment. We leave it as an exercise for the reader to prove that, based on (1.46) and barring sampling variations, the percentage Q of deceased subjects from the treatment group who would have recovered had they not taken the treatment precisely equals the percentage Q' of surviving subjects in the nontreatment group who will die if given treatment. Q' is hypothetical, Q' is unquestionably testable.

Having sketched the empirical interpretation of counterfactuals, our next step in this introductory chapter is the question of representation: What knowledge is required to answer questions about counterfactuals? And how should this knowledge be formulated so that counterfactual queries can be answered quickly and reliably? That such representation exists is evident by the swiftness and consistency with which people distinguish plausible from implausible counterfactual statements. Most people would agree that President Clinton's place in history would be different had he not met Monica Lewinsky, but only a few would assert that his place in history would change had he not eaten breakfast yesterday. In the cognitive sciences, such consistency of opinion is as close as one can get to a proof that an effective machinery for representing and manipulating counterfactuals resides someplace in the human mind. What then are the building blocks of that machinery?

A straightforward representational scheme would (i) store counterfactual knowledge in the form of counterfactual premises and (ii) derive answers to counterfactual queries using some logical rules of inference capable of taking us from premises to conclusions. This approach has indeed been taken by the philosophers Robert Stalnaker (1968) and David Lewis (1973a,b), who constructed logics of counterfactuals using closest-world

<sup>&</sup>lt;sup>23</sup> For example, if *Q* equals 100% (i.e., all those who took the treatment and died would have recovered had they not taken the treatment), then all surviving subjects from the nontreatment group will die if given treatment (again, barring sampling variations). Such exercises will become routine when we develop the mathematical machinery for analyzing probabilities of causes (see Chapter 9, Theorem 9.2.12, equations (9.11)–(9.12)).



**Figure 1.6** (a) A causal Bayesian network that represents the distribution of (1.47). (b) A causal diagram representing the process generating the distribution in (a), according to model 1. (c) Same, according to model 2. (Both  $U_1$  and  $U_2$  are unobserved.)

semantics (i.e., "B would be true if it were A" just in case B is true in the closest possible world (to ours) in which A is true). However, the closest-world semantics still leaves two questions unanswered. (1) What choice of distance measure would make counterfactual reasoning compatible with ordinary conceptions of cause and effect? (2) What mental representation of interworld distances would render the computation of counterfactuals manageable and practical (for both humans and machines)? These two questions are answered by the structural model approach expanded in Chapter 7.

An approach similar to Lewis's (though somewhat less formal) has been pursued by statisticians in the potential-outcome framework (Rubin 1974; Robins 1986; Holland 1988). Here, substantive knowledge is expressed in terms of probabilistic relationships (e.g., independence) among counterfactual variables and then used in the estimation of causal effects. The question of representation shifts from the closest-world to the potential-outcome approach: How are probabilistic relationships among counterfactuals stored or inferred in the investigator's mind? In Chapter 7 (see also Section 3.6.3) we provide an analysis of the closest-world and potential-outcome approaches and compare them to the structural model approach, to be outlined next, in which counterfactuals are derived from (and in fact defined by) a functional causal model (equation (1.40)).

In order to see the connection between counterfactuals and structural equations, we should first examine why the information encoded in a Bayesian network, even in its causal interpretation, is insufficient to answer counterfactual queries. Consider again our example of the controlled randomized experiment (equation (1.46)), which corresponds to an edgeless Bayesian network (Figure 1.6(a)) with two independent binary variables and a joint probability:

$$P(y, x) = 0.25$$
 for all x and y. (1.47)

We now present two functional models, each generating the joint probability of (1.47) yet each giving a different value to the quantity of interest, Q = the probability that a subject who died under treatment (x = 1, y = 1) would have recovered (y = 0) had he or she not been treated (x = 0).

Model 1 (Figure 1.6(b))  
Let  

$$x = u_1$$
,  
 $y = u_2$ .

where  $U_1$  and  $U_2$  are two independent binary variables with  $P(u_1 = 1) = P(u_2 = 1) = \frac{1}{2}$  (e.g., random coins).

linearity, normality, and error independence are sufficient for computing all counterfactual queries in the model of Figure 1.5 (see Section 7.2.1). In Chapter 9, we will present conditions under which counterfactual queries concerning probability of causation can be inferred from data when  $f_i$  and P(u) are unknown, and only general features (e.g., monotonicity) of these entities are assumed. Likewise, Chapter 8 (Section 8.3) will present methods of *bounding* probabilities of counterfactuals when only stochastic models are available.

The preceding considerations further imply that the three tasks listed in the beginning of this section – prediction, intervention, and counterfactuals – form a natural hierarchy of causal reasoning tasks, with increasing levels of refinement and increasing demands on the knowledge required for accomplishing these tasks. Prediction is the simplest of the three, requiring only a specification of a joint distribution function. The analysis of interventions requires a causal structure in addition to a joint distribution. Finally, processing counterfactuals is the hardest task because it requires some information about the functional relationships and/or the distribution of the omitted factors.

This hierarchy also defines a natural partitioning of the chapters in this book. Chapter 2 will deal primarily with the probabilistic aspects of causal Bayesian networks (though the underlying causal structure will serve as a conceptual guide). Chapters 3–6 will deal exclusively with the interventional aspects of causal models, including the identification of causal effects, the clarification of structural equation models, and the relationships between confounding and collapsibility. Chapters 7–10 will deal with counterfactual analysis, including axiomatic foundation, applications to policy analysis, the bounding of counterfactual queries, the identification of probabilities of causes, and the explication of single-event causation.

I wish the reader a smooth and rewarding journey through these chapters. But first, an important stop for terminological distinctions.

#### 1.5 CAUSAL VERSUS STATISTICAL TERMINOLOGY

This section defines fundamental terms and concepts that will be used throughout this book. These definitions may not agree with those given in standard sources, so it is important to refer to this section in case of doubts regarding the interpretation of these terms.

A *probabilistic parameter* is any quantity that is defined in terms<sup>25</sup> of a joint probability function. Examples are the quantities defined in Sections 1.1 and 1.2.

A *statistical parameter* is any quantity that is defined in terms of a joint probability distribution of observed variables, making no assumption whatsoever regarding the existence or nonexistence of unobserved variables.

```
Examples: the conditional expectation E(Y | x), the regression coefficient r_{YX}, the value of the density function at y = 0, x = 1.
```

A *causal parameter* is any quantity that is defined in terms of a causal model (as in (1.40)) and is not a statistical parameter.

A quantity Q is said to be defined in terms of an object of class C if Q can be computed uniquely from the description of any object in class C (i.e., if Q is defined by a functional mapping from C to the domain of Q).

**Examples:** the coefficients  $a_{ik}$  in (1.41),

whether  $X_0$  has influence on  $X_3$  for some u,

the expected value of Y under the intervention do(X = 0),

the number of parents of variable  $X_7$ .

**Remark:** The exclusion of unmeasured variables from the definition of statistical parameters is devised to prevent one from hiding causal assumptions under the guise of latent variables. Such constructions, if permitted, would qualify any quantity as statistical and would thus obscure the important distinction between quantities that can be estimated from statistical data alone, and those that require additional assumptions beyond the data.

A *statistical assumption* is any constraint on a joint distribution of observed variable; for example, that *f* is multivariate normal or that *P* is Markov relative to a given DAG *D*.

A *causal assumption* is any constraint on a causal model that cannot be realized by imposing statistical assumptions; for example, that  $f_i$  is linear, that  $U_i$  and  $U_j$  (unobserved) are uncorrelated, or that  $x_3$  does not appear in  $f_4(pa_4, u_4)$ . Causal assumptions may or may not have statistical implications. In the former case we say that the assumption is "testable" or "falsifiable." Often, though not always, causal assumptions can be falsified from experimental studies, in which case we say that they are "experimentally testable." For example, the assumption that X has no effect on E(Y) in model 2 of Figure 1.6 is empirically testable, but the assumption that X may cure a given subject in the population is not.

**Remark:** The distinction between causal and statistical parameters is crisp and fundamental – the two do not mix. Causal parameters cannot be discerned from statistical parameters unless causal assumptions are invoked. The formulation and simplification of these assumptions will occupy a major part of this book.

**Remark:** Temporal precedence among variables may furnish some information about (the absence of) causal relationships – a later event cannot be the cause of an earlier event. Temporally indexed distributions such as  $P(y_t \mid y_{t-1}, x_t)$ , t = 1,..., which are used routinely in economic analysis, may therefore be regarded as borderline cases between statistical and causal models. We shall nevertheless classify those models as statistical because the great majority of policy-related questions *cannot* be discerned from such distributions, given our commitment to making no assumption regarding the presence or absence of unmeasured variables. Consequently, econometric concepts such as "Granger causality" (Granger 1969) and "strong exogeneity" (Engle et al. 1983) will be classified as statistical rather than causal.<sup>26</sup>

**Remark:** The terms "theoretical" and "structural" are often used interchangeably with "causal"; we will use the latter two, keeping in mind that some structural models may not be causal (see Section 7.2.5).

Caution must also be exercised in labeling as a "data-generating model" the probabilistic sequence  $P(y_t|y_{t-1}x_t)$ ,  $t=1,\ldots$  (e.g., Davidson and MacKinnon 1993, p. 53; Hendry 1995). Such sequences are statistical in nature and, unless causal assumptions of the type developed in Chapter 2 (see Definitions 2.4.1 and 2.7.4) are invoked, they cannot be applied to policy-evaluation tasks.

## Causal versus Statistical Concepts

The demarcation line between causal and statistical parameters extends as well to general concepts and will be supported by terminological distinction. Examples of *statistical* concepts are: correlation, regression, conditional independence, association, likelihood, collapsibility, risk ratio, odds ratio, propensity score, Granger's causality, and so on. Examples of *causal* concepts are: randomization, influence, effect, confounding, exogeneity, ignorability, disturbance (e.g., (1.40)), spurious correlation, path coefficients, instrumental variables, intervention, explanation, and so on. The purpose of this demarcation line is not to exclude causal concepts from the province of statistical analysis but, rather, to encourage investigators to treat nonstatistical concepts with the proper set of tools.

Some readers may be surprised by the idea that textbook concepts such as randomization, confounding, spurious correlation, and effects are nonstatistical. Others may be shocked at the idea that controversial concepts such as exogeneity, confounding, and counterfactuals *can* be defined in terms of causal models. This book is written with these readers in mind, and the coming pages will demonstrate that the distinctions just made between causal and statistical concepts are essential for clarifying both.

## Two Mental Barriers to Causal Analysis

The sharp distinction between statistical and causal concepts can be translated into a useful principle: behind every causal claim there must lie some causal assumption that is not discernable from the joint distribution and, hence, not testable in observational studies. Such assumptions are usually provided by humans, resting on expert *judgment*. Thus, the way humans organize and communicate experiential knowledge becomes an integral part of the study, for it determines the veracity of the judgments experts are requested to articulate.

Another ramification of this causal–statistical distinction is that any mathematical approach to causal analysis must acquire *new notation*. The vocabulary of probability calculus, with its powerful operators of expectation, conditionalization, and marginalization, is defined strictly in terms of distribution functions and is therefore insufficient for expressing causal assumptions or causal claims. To illustrate, the syntax of probability calculus does not permit us to express the simple fact that "symptoms do not cause diseases," let alone draw mathematical conclusions from such facts. All we can say is that two events are dependent – meaning that if we find one, we can expect to encounter the other, but we cannot distinguish statistical dependence, quantified by the conditional probability  $P(disease \mid symptom)$ , from causal dependence, for which we have no expression in standard probability calculus.

The preceding two requirements: (1) to commence causal analysis with untested, judgmental assumptions, and (2) to extend the syntax of probability calculus, constitute the two main obstacles to the acceptance of causal analysis among professionals with traditional training in statistics (Pearl 2003c, also sections 11.1.1 and 11.6.4). This book helps overcome the two barriers through an effective and friendly notational system based on symbiosis of graphical and algebraic approaches.

# A Theory of Inferred Causation

I would rather discover one causal law than be King of Persia.

Democritus (460–370 B.C.)

## **Preface**

The possibility of learning causal relationships from raw data has been on philosophers' dream lists since the time of Hume (1711–1776). That possibility entered the realm of formal treatment and feasible computation in the mid-1980s, when the mathematical relationships between graphs and probabilistic dependencies came to light. The approach described herein is an outgrowth of Rebane and Pearl (1987) and Pearl (1988b, Chap. 8), which describes how causal relationships can be inferred from nontemporal statistical data if one makes certain assumptions about the underlying process of data generation (e.g., that it has a tree structure). The prospect of inferring causal relationships from weaker structural assumptions (e.g., general directed acyclic graphs) has motivated parallel research efforts at three universities: UCLA, Carnegie Mellon University (CMU), and Stanford. The UCLA and CMU teams pursued an approach based on searching the data for patterns of conditional independencies that reveal fragments of the underlying structure and then piecing those fragments together to form a coherent causal model (or a set of such models). The Stanford group pursued a Bayesian approach, where data are used to update prior probabilities assigned to candidate causal structures (Cooper and Herskovits 1991). The UCLA and CMU efforts have led to similar theories and almost identical discovery algorithms, which were implemented in the TETRAD II program (Spirtes et al. 1993). The Bayesian approach has since been pursued by a number of research teams (Singh and Valtorta 1995; Heckerman et al. 1994) and now serves as the basis for several graph-based learning methods (Jordan 1998). This chapter describes the approach pursued by Tom Verma and me in the period 1988-1992, and it briefly summarizes related extensions, refinements, and improvements that have been advanced by the CMU team and others. Some of the philosophical rationale behind this development, primarily the assumption of minimality, are implicit in the Bayesian approach as well (Section 2.9.1).

The basic idea of automating the discovery of causes – and the specific implementation of this idea in computer programs – came under fierce debate in a number of forums (Cartwright 1995a; Humphreys and Freedman 1996; Cartwright 1999; Korb and Wallace 1997; McKim and Turner 1997; Robins and Wasserman 1999). Selected aspects of this debate will be addressed in the discussion section at the end of this chapter (Section 2.9.1).

Acknowledging that statistical associations do not *logically* imply causation, this chapter asks whether weaker relationships exist between the two. In particular, we ask:

- 1. What clues prompt people to perceive causal relationships in uncontrolled observations?
- 2. What assumptions would allow us to infer causal models from these clues?
- 3. Would the models inferred tell us anything useful about the causal mechanisms that underly the observations?

In Section 2.2 we define the notions of causal models and causal structures and then describe the task of causal discovery as an inductive game that scientists play against Nature. In Section 2.3 we formalize the inductive game by introducing "minimal model" semantics - the semantical version of Occam's razor - and exemplify how, contrary to common folklore, causal relationships can be distinguished from spurious covariations following this standard norm of inductive reasoning. Section 2.4 identifies a condition, called stability (or faithfulness), under which effective algorithms exist that uncover structures of casual influences as defined here. One such algorithm (called IC), introduced in Section 2.5, uncovers the set of all causal models compatible with the data, assuming all variables are observed. Another algorithm (IC\*), described in Section 2.6, is shown to uncover many (though not all) valid causal relationships when some variables are not observable. In Section 2.7 we extract from the IC\* algorithm the essential conditions under which causal influences are inferred, and we offer these as independent definitions of genuine influences and spurious associations, with and without temporal information. Section 2.8 offers an explanation for the puzzling yet universal agreement between the temporal and statistical aspects of causation. Finally, Section 2.9 summarizes the claims made in this chapter, re-explicates the assumptions that lead to these claims, and offers new justifications of these assumption in light of ongoing debates.

## 2.1 INTRODUCTION - THE BASIC INTUITIONS

An autonomous intelligent system attempting to build a workable model of its environment cannot rely exclusively on preprogrammed causal knowledge; rather, it must be able to translate direct observations to cause-and-effect relationships. However, given that statistical analysis is driven by covariation, not causation, and assuming that the bulk of human knowledge derives from passive observations, we must still identify the clues that prompt people to perceive causal relationships in the data. We must also find a computational model that emulates this perception.

Temporal precedence is normally assumed to be essential for defining causation, and it is undoubtedly one of the most important clues that people use to distinguish causal from other types of associations. Accordingly, most theories of causation invoke an explicit requirement that a cause precede its effect in time (Reichenbach 1956; Good 1961; Suppes 1970; Shoham 1988). Yet temporal information alone cannot distinguish genuine causation from spurious associations caused by unknown factors – the barometer falls before it rains yet does not cause the rain. In fact, the statistical and philosophical literature has adamantly warned analysts that, unless one knows in advance all causally

the observables, but it hides the underlying causal model as well as the causal structure. We investigate the feasibility of recovering the topology D of the DAG from features of the probability distribution  $P_{[O]}$ .<sup>2</sup>

#### 2.3 MODEL PREFERENCE (OCCAM'S RAZOR)

In principle, since V is unknown, there is an unbounded number of models that would fit a given distribution, each invoking a different set of "hidden" variables and each connecting the observed variables through different causal relationships. Therefore, with no restriction on the type of models considered, the scientist is unable to make any meaningful assertions about the structure underlying the phenomena. For example, every probability distribution  $P_{[O]}$  can be generated by a structure in which no observed variable is a cause of another but instead all variables are consequences of one latent common cause, U. Likewise, assuming V = O but lacking temporal information, the scientist can never rule out the possibility that the underlying structure is a complete, acyclic, and arbitrarily ordered graph — a structure that (with the right choice of parameters) can *mimic* the behavior of any model, regardless of the variable ordering. However, following standard norms of scientific induction, it is reasonable to rule out any theory for which we find a simpler, less elaborate theory that is equally consistent with the data (see Definition 2.3.5). Theories that survive this selection process are called *minimal*. With this notion, we can construct our (preliminary) definition of inferred causation as follows.

## **Definition 2.3.1 (Inferred Causation (Preliminary))**

A variable X is said to have a causal influence on a variable Y if a directed path from X to Y exists in every minimal structure consistent with the data.

Here we equate a causal structure with a scientific theory, since both contain a set of free parameters that can be adjusted to fit the data. We regard Definition 2.3.1 as preliminary because it assumes that all variables are observed. The next few definitions generalize the concept of minimality to structures with unobserved variables.

#### **Definition 2.3.2 (Latent Structure)**

A latent structure is a pair  $L = \langle D, O \rangle$ , where D is a causal structure over V and where  $O \subseteq V$  is a set of observed variables.

## **Definition 2.3.3 (Structure Preference)**

One latent structure  $L = \langle D, O \rangle$  is preferred to another  $L' = \langle D', O \rangle$  (written  $L \leq L'$ ) if and only if D' can mimic D over O – that is, if and only if for every  $\Theta_D$  there exists a

<sup>&</sup>lt;sup>2</sup> This formulation invokes several idealizations of the actual task of scientific discovery. It assumes, for example, that the scientist obtains the distribution directly, rather than events sampled from the distribution. Additionally, we assume that the observed variables actually appear in the original causal model and are not some aggregate thereof. Aggregation might result in feedback loops, which we do not discuss in this chapter.

<sup>&</sup>lt;sup>3</sup> This can be realized by letting U have as many states as O, assigning to U the prior distribution P(u) = P(o(u)) (where o(u) is the cell of O corresponding to state u), and letting each observed variable  $O_i$  take on its corresponding value in o(u).

 $\Theta'_{D'}$  such that  $P_{[O]}(\langle D', \Theta'_{D'} \rangle) = P_{[O]}(\langle D, \Theta_D \rangle)$ . Two latent structures are equivalent, written  $L' \equiv L$ , if and only if  $L \preceq L'$  and  $L \succeq L'$ .<sup>4</sup>

Note that the preference for simplicity imposed by Definition 2.3.3 is gauged by the expressive power of a structure, not by its syntactic description. For example, one latent structure  $L_1$  may invoke many more parameters than  $L_2$  and still be preferred if  $L_2$  can accommodate a richer set of probability distributions over the observables. One reason scientists prefer simpler theories is that such theories are more constraining and thus more falsifiable; they provide the scientist with less opportunities to overfit the data "hindsightedly" and therefore command greater credibility if a fit is found (Popper 1959; Pearl 1978; Blumer et al. 1987).

We also note that the set of independencies entailed by a causal structure imposes limits on its expressive power, that is, its power to mimic other structures. Indeed,  $L_1$  cannot be preferred to  $L_2$  if there is even one observable dependency that is permitted by  $L_1$  and forbidden by  $L_2$ . Thus, tests for preference and equivalence can sometimes be reduced to tests of induced dependencies, which in turn can be determined directly from the topology of the DAGs without ever concerning ourselves with the set of parameters. This is the case in the absence of hidden variables (see Theorem 1.2.8) but does not hold generally in all latent structures. Verma and Pearl (1990) showed that some latent structures impose numerical rather than independence constraints on the observed distribution (see, e.g., Section 8.4, equations (8.21)–(8.23)); this makes the task of verifying model preference complicated but does still permit us to extend the semantical definition of inferred causation (Definition 2.3.1) to latent structures.

## **Definition 2.3.4 (Minimality)**

A latent structure L is minimal with respect to a class  $\mathcal{L}$  of latent structures if and only if there is no member of  $\mathcal{L}$  that is strictly preferred to L – that is, if and only if for every  $L' \in \mathcal{L}$  we have  $L \equiv L'$  whenever  $L' \prec L$ .

## **Definition 2.3.5 (Consistency)**

A latent structure L = |D,O| is consistent with a distribution  $\hat{P}$  over O if D can accommodate some model that generates  $\hat{P}$  – that is, if there exists a parameterization  $\Theta_D$  such that  $P_{[O]}(\langle D,\Theta_D\rangle)=\hat{P}$ .

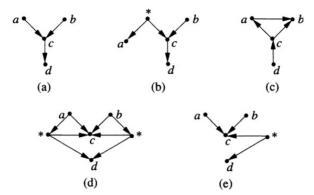
Clearly, a necessary (and sometimes sufficient) condition for L to be consistent with  $\hat{P}$  is that L can account for all the dependencies embodied in  $\hat{P}$ 

#### **Definition 2.3.6 (Inferred Causation)**

Given  $\hat{P}$ , a variable C has a causal influence on variable E if and only if there exists a directed path from C to E in every minimal latent structure consistent with  $\hat{P}$ .

We view this definition as normative because it is based on one of the least disputed norms of scientific investigation: Occam's razor in its semantical casting. However, as with any

<sup>&</sup>lt;sup>4</sup> We use the succinct term "preferred to" to mean "preferred or equivalent to," a relation that has also been named "a submodel of."



**Figure 2.1** Causal structures illustrating the minimality of (a) and (b) and the justification for inferring the relationship  $c \to d$ . The node (\*) represents a hidden variable with any number of states.

scientific inquiry, we make no claims that this definition is guaranteed to always identify stable physical mechanisms in Nature. It identifies the mechanisms we can plausibly infer from nonexperimental data; moreover, it guarantees that any alternative mechanism will be less trustworthy than the one inferred because the alternative would require more contrived, hindsighted adjustment of parameters (i.e., functions) to fit the data.

As an example of a causal relation that is identified by Definition 2.3.6, imagine that observations taken over four variables  $\{a, b, c, d\}$  reveal two independencies: "a is independent of b" and "d is independent of  $\{a, b\}$  given c." Assume further that the data reveals no other independence besides those that logically follow from these two. This dependence pattern would be typical, for example, of the following variables: a = havinga cold, b = having hav fever, c = having to sneeze, d = having to wipe one's nose. It is not hard to see that structures (a) and (b) in Figure 2.1 are minimal, for they entail the observed independencies and none other.<sup>5</sup> Furthermore, any structure that explains the observed dependence between c and d by an arrow from d to c, or by a hidden common cause (\*) between the two, cannot be minimal, because any such structure would be able to "out-mimic" the one shown in Figure 2.1(a) (or the one in Figure 2.1(b)), which reflects all observed independencies. For example, the structure of Figure 2.1(c), unlike that of Figure 2.1(a), accommodates distributions with arbitrary relations between a and b. Similarly, Figure 2.1(d) is not minimal because it fails to impose the conditional independence between d and  $\{a, b\}$  given c and will therefore accommodate distributions in which d and  $\{a, b\}$  are dependent given c. In contrast, Figure 2.1(e) is not consistent with the data, since it imposes an unobserved marginal independence between  $\{a, b\}$ and d.

This example (taken from Pearl and Verma 1991) illustrates a remarkable connection between causality and probability: certain patterns of probabilistic dependencies (in our case, all dependencies except  $(a \perp \!\!\!\perp b)$  and  $(d \perp \!\!\!\perp \{a,b\} \mid c)$ ) imply unambiguous *causal* dependencies (in our case,  $c \rightarrow d$ ) without making any assumption about the presence

<sup>&</sup>lt;sup>5</sup> To verify that (a) and (b) are equivalent, we note that (b) can mimic (a) if we let the link *a* ← \* impose equality between the two variables. Conversely, (a) can mimic (b), since it is capable of generating every distribution that possesses the independencies entailed by (b). (For theory and methods of "reading off" conditional independencies from graphs, see Section 1.2.3 or Pearl 1988b.)

or absence of latent variables.<sup>6</sup> The only assumption invoked in this implication is minimality – models that overfit the data are ruled out.

#### 2.4 STABLE DISTRIBUTIONS

Although the minimality principle is sufficient for forming a normative theory of inferred causation, it does not guarantee that the structure of the actual data-generating model would be minimal, or that the search through the vast space of minimal structures would be computationally practical. Some structures may admit peculiar parameterizations that would render them indistinguishable from many other minimal models that have totally disparate structures. For example, consider a binary variable C that takes the value 1 whenever the outcomes of two fair coins (A and B) are the same and takes the value 0 otherwise. In the trivariate distribution generated by this parameterization, each pair of variables is marginally independent yet is dependent conditional on the third variable. Such a dependence pattern may in fact be generated by three minimal causal structures, each depicting one of the variables as causally dependent on the other two, but there is no way to decide among the three. In order to rule out such "pathological" parameterizations, we impose a restriction on the distribution called *stability*, also known as DAG-isomorphism or perfect-mapness (Pearl 1988b, p. 128) and faithfulness (Spirtes et al. 1993). This restriction conveys the assumption that all the independencies embedded in P are stable; that is, they are entailed by the structure of the model D and hence remain invariant to any change in the parameters  $\Theta_D$ . In our example, only the correct structure (namely,  $A \to C \leftarrow B$ ) will retain its independence pattern in the face of changing parameterizations – say, when the coins become slightly biased.

## **Definition 2.4.1 (Stability)**

Let I(P) denote the set of all conditional independence relationships embodied in P. A causal model  $M = \langle D, \Theta_D \rangle$  generates a stable distribution if and only if  $P(\langle D, \Theta_D \rangle)$  contains no extraneous independences – that is, if and only if  $I(P(\langle D, \Theta_D \rangle)) \subseteq I(P(\langle D, \Theta_D \rangle))$  for any set of parameters  $\Theta_D$ .

The stability condition states that, as we vary the parameters from  $\Theta$  to  $\Theta'$ , no independence in P can be destroyed; hence the name "stability." Succinctly, P is a stable distribution of M if it "maps" the structure D of M, that is,  $(X \perp \!\!\!\perp Y \mid Z)_P \iff (X \perp \!\!\!\perp Y \mid Z)_D$  for any three sets of variables X, Y, and Z (see Theorem 1.2.5).

The relationship between minimality and stability can be illustrated using the following analogy. Suppose we see a picture of a chair and that we need to decide between two theories as follows.

- $T_1$ : The object in the picture is a chair.
- T<sub>2</sub>: The object in the picture is either a chair or two chairs positioned such that one hides the other.

<sup>&</sup>lt;sup>6</sup> Standard probabilistic definitions of causality (e.g., Suppes 1970; Eells 1991) invariably require knowledge of all relevant factors that may influence the observed variables (see Section 7.5.3).

Our preference for  $T_1$  over  $T_2$  can be justified on two principles, one based on minimality and the other on stability. The minimality principle argues that  $T_1$  is preferred to  $T_2$  because the set of scenes composed of single objects is a proper subset of scenes composed of two or fewer objects and, unless we have evidence to the contrary, we should prefer the more specific theory. The stability principle rules out  $T_2$  a priori, arguing that it would be rather unlikely for two objects to align themselves so as to have one perfectly hide the other. Such an alignment would be unstable relative to slight changes in environmental conditions or viewing angle.

The analogy with independencies is clear. Some independencies are *structural*, that is, they would persist for every functional—distributional parameterization of the graph. Others are sensitive to the precise numerical values of the functions and distributions. For example, in the structure  $Z \leftarrow X \rightarrow Y$ , which stands for the relations

$$z = f_1(x, u_1), \quad y = f_2(x, u_2),$$
 (2.1)

the variables Z and Y will be independent, conditional on X, for all functions  $f_1$  and  $f_2$ . In contrast, if we add an arrow  $Z \rightarrow Y$  to the structure and use a linear model

$$z = \gamma x + u_1, \quad y = \alpha x + \beta z + u_2, \tag{2.2}$$

with  $\alpha = -\beta \gamma$ , then Y and X will be independent. However, the independence between Y and X is unstable because it disappears as soon as the equality  $\alpha = -\beta \gamma$  is violated. The stability assumption presumes that this type of independence is unlikely to occur in the data, that all independencies are structural.

To further illustrate the relations between stability and minimality, consider the causal structure depicted in Figure 2.1(c). The minimality principle rejects this structure on the ground that it fits a broader set of distributions than those fitted by structure (a). The stability principle rejects this structure on the ground that, in order to fit the data (specifically, the independence  $(a \perp \!\!\!\perp b)$ ), the association produced by the arrow  $a \rightarrow b$  must cancel precisely the one produced by the path  $a \leftarrow c \rightarrow b$ . Such precise cancelation cannot be stable, for it cannot be sustained for all functions connecting variables a, b, and c. In structure (a), by contrast, the independence  $(a \perp \!\!\!\perp b)$  is stable.

#### 2.5 RECOVERING DAG STRUCTURES

With the added assumption of stability, every distribution has a unique minimal causal structure (up to d-separation equivalence), as long as there are no hidden variables. This uniqueness follows from Theorem 1.2.8, which states that two causal structures are equivalent (i.e., they can mimic each other) if and only if they relay the same dependency information – namely, they have the same skeleton and same set of v-structures.

In the absence of unmeasured variables, the search for the minimal model then boils down to reconstructing the structure of a DAG D from queries about conditional independencies, assuming that those independencies reflect d-separation conditions in some undisclosed underlying DAG  $D_0$ . Naturally, since  $D_0$  may have equivalent structures, the reconstructed DAG will not be unique, and the best we can do is to find a graphical representation for the equivalence class of  $D_0$ . Such graphical representation was introduced in Verma and Pearl (1990) under the name pattern. A pattern is a partially directed

#### **Definition 2.6.1 (Projection)**

A latent structure  $L_{[O]} = \langle D_{[O]}, O \rangle$  is a projection of another latent structure L if and only if:

- 1. every unobservable variable of  $D_{[O]}$  is a parentless common cause of exactly two nonadjacent observable variables; and
- 2. for every stable distribution P generated by L, there exists a stable distribution P' generated by  $L_{[O]}$  such that  $I(P_{[O]}) = I(P'_{[O]})$ .

#### **Theorem 2.6.2** (Verma 1993)

Any latent structure has at least one projection.

It is convenient to represent projections using a bidirectional graph with only the observed variables as vertices (i.e., leaving the hidden variables implicit). Each bidirected link in such a graph represents a common hidden cause of the variables corresponding to the link's endpoints.

Theorem 2.6.2 renders our definition of inferred causation (Definition 2.3.6) operational; it can be shown (Verma 1993) that the existence of a certain link in a distinguished projection of any minimal model of  $\hat{P}$  must indicate the existence of a causal path in every minimal model of  $\hat{P}$ . Thus, our search reduces to finding the distinguished projection of any minimal model of  $\hat{P}$  and identifying the appropriate links. Remarkably, these links can be identified by a simple variant of the IC algorithm, here called IC\*, that takes a stable distribution  $\hat{P}$  and returns a *marked* pattern, which is a partially directed acyclic graph that contains four types of edges:

- 1. a marked arrow  $a \xrightarrow{*} b$ , signifying a directed path from a to b in the underlying model;
- 2. an unmarked arrow  $a \rightarrow b$ , signifying either a directed path from a to b or a latent common cause  $a \leftarrow L \rightarrow b$  in the underlying model;
- 3. a bidirected edge  $a \longleftrightarrow b$ , signifying a latent common cause  $a \leftarrow L \rightarrow b$  in the underlying model; and
- 4. an undirected edge a b, standing for either  $a \leftarrow b$  or  $a \rightarrow b$  or  $a \leftarrow L \rightarrow b$  in the underlying model.<sup>8</sup>

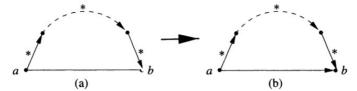
## IC\* Algorithm (Inductive Causation with Latent Variables)

**Input:**  $\hat{P}$ , a stable distribution (with respect to some latent structure).

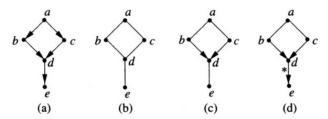
**Output:** core  $(\hat{P})$ , a marked pattern.

1. For each pair of variables a and b, search for a set  $S_{ab}$  such that a and b are independent in  $\hat{P}$ , conditioned on  $S_{ab}$ .

<sup>8</sup> Spirtes et al. (1993) used a o→b to represent uncertainty about the arrowhead at node a. Several errors in the original proof of IC\* were pointed out to us by Peter Spirtes and were corrected in Verma (1993). Alternative proofs of correctness, as well as refinements in the algorithm, are given in Spirtes et al. (1993).



**Figure 2.2** Illustration of  $R_2$  in step 3 of the IC\* algorithm.



**Figure 2.3** Graphs constructed by the IC\* algorithm. (a) Underlying structure. (b) After step 1. (c) After step 2. (d) Output of IC\*.

If there is no such  $S_{ab}$ , place an undirected link between the two variables, a-b.

2. For each pair of nonadjacent variables a and b with a common neighbor c, check if  $c \in S_{ab}$ .

If it is, then continue.

If it is not, then add arrowheads pointing at c (i.e.,  $a \rightarrow c \leftarrow b$ ).

- 3. In the partially directed graph that results, add (recursively) as many arrowheads as possible, and mark as many edges as possible, according to the following two rules:
  - $R_1$ : For each pair of nonadjacent nodes a and b with a common neighbor c, if the link between a and c has an arrowhead into c and if the link between c and b has no arrowhead into c, then add an arrowhead on the link between c and b pointing at b and mark that link to obtain  $c \stackrel{*}{\longrightarrow} b$ .
  - R<sub>2</sub>: If a and b are adjacent and there is a directed path (composed strictly of marked links) from a to b (as in Figure 2.2), then add an arrowhead pointing toward b on the link between a and b.

Steps 1 and 2 of IC\* are identical to those of IC, but the rules in step 3 are different; they do not orient edges but rather add arrowheads to the individual endpoints of the edges, thus accommodating bidirectional edges.

Figure 2.3 illustrates the operation of the IC\* algorithm on the sprinkler example of Figure 1.2 (shown schematically in Figure 2.3(a)).

1. The conditional independencies entailed by this structure can be read off using the *d*-separation criterion (Definition 1.2.3), and the smallest conditioning sets corresponding to these independencies are given by  $S_{ad} = \{b, c\}$ ,  $S_{ae} = \{d\}$ ,

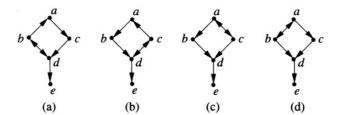


Figure 2.4 Latent structures equivalent to those of Figure 2.3(a).

 $S_{bc} = \{a\}$ ,  $S_{be} = \{d\}$ , and  $S_{ce} = \{d\}$ . Thus, step 1 of IC\* yields the undirected graph of Figure 2.3(b).

- 2. The triplet (b, d, c) is the only one that satisfies the condition of step 2, since d is not in  $S_{bc}$ . Accordingly, we obtain the partially directed graph of Figure 2.3(c).
- 3. Rule  $R_1$  of step 3 is applicable to the triplet (b, d, e) (and to (c, d, e)), since b and e are nonadjacent and there is an arrowhead at d from b but not from e. We therefore add an arrowhead at e, and mark the link, to obtain Figure 2.3(d). This is also the final output of IC\*, because  $R_1$  and  $R_2$  are no longer applicable.

The absence of arrowheads on a-b and a-c, and the absence of markings on  $b \to d$  and  $c \to d$ , correctly represent the ambiguities presented by  $\hat{P}$ . Indeed, each of the latent structures shown in Figure 2.4 is observationally equivalent to that of Figure 2.3(a). Marking the link  $d \to e$  in Figure 2.3(d) advertises the existence of a directed link  $d \to e$  in each and every latent structure that is independence-equivalent to the one in Figure 2.3(a).

## 2.7 LOCAL CRITERIA FOR INFERRING CAUSAL RELATIONS

The IC\* algorithm takes a distribution  $\hat{P}$  and outputs a partially directed graph. Some of the links are marked unidirectional (denoting genuine causation), some are unmarked unidirectional (denoting potential causation), some are bidirectional (denoting spurious association), and some are undirected (denoting relationships that remain undetermined). The conditions that give rise to these labelings can be taken as definitions for the various kinds of causal relationships. In this section we present explicit definitions of potential and genuine causation as they emerge from the IC\* algorithm. Note that, in all these definitions, the criterion for causation between two variables (X and Y) will require that a third variable Z exhibit a specific pattern of dependency with X and Y. This is not surprising, since the essence of causal claims is to stipulate the behavior of X and Y under the influence of a third variable, one that corresponds to an external control of X (or Y) – as echoed in the paradigm of "no causation without manipulation" (Holland 1986). The difference is only that the variable Z, acting as a virtual control, must be identified within the data itself, as if Nature had performed the experiment. The IC\* algorithm can be regarded as offering a systematic way of searching for variables Z that qualify as virtual controls, given the assumption of stability.

#### **Definition 2.7.1 (Potential Cause)**

A variable X has a potential causal influence on another variable Y (that is inferable from  $\hat{P}$ ) if the following conditions hold.

- 1. X and Y are dependent in every context.
- 2. There exists a variable Z and a context S such that
  - (i) X and Z are independent given S (i.e.,  $X \perp \!\!\! \perp Z \mid S$ ) and
  - (ii) Z and Y are dependent given S (i.e.,  $Z \perp \!\!\! \perp Y \mid S$ ).

By "context" we mean a set of variables tied to specific values. In Figure 2.3(a), for example, variable b qualifies as a potential cause of d by virtue of variable Z = c being dependent on d and independent of b in context S = a. Likewise, c qualifies as a potential cause of d (with Z = b and S = a). Neither b nor c qualifies as a genuine cause of d, because this pattern of dependencies is also compatible with a latent common cause, shown as bidirected arcs in Figures 2.4(a)–(b). However, Definition 2.7.1 disqualifies d as a cause of b (or c), and this leads to the classification of d as a genuine cause of e, as formulated in Definition 2.7.2. Note that Definition 2.7.1 precludes a variable X from being a potential cause of itself or of any other variable that functionally determines X.

#### **Definition 2.7.2 (Genuine Cause)**

A variable X has a genuine causal influence on another variable Y if there exists a variable Z such that either:

- 1. X and Y are dependent in any context and there exists a context S satisfying
  - (i) Z is a potential cause of X (per Definition 2.7.1),
  - (ii) Z and Y are dependent given S (i.e.,  $Z \not\perp \!\!\! \perp Y \mid S$ ), and
  - (iii) Z and Y are independent given  $S \cup X$  (i.e.,  $Z \perp\!\!\!\perp Y \mid S \cup X$ ); or
- 2. X and Y are in the transitive closure of the relation defined in criterion 1.

Conditions (i)–(iii) are illustrated in Figure 2.3(a) with X = d, Y = e, Z = b, and  $S = \emptyset$ . The destruction of the dependence between b and e through conditioning on d cannot be attributed to spurious association between d and e; genuine causal influence is the only explanation, as shown in the structures of Figure 2.4.

#### **Definition 2.7.3 (Spurious Association)**

Two variables X and Y are spuriously associated if they are dependent in some context and there exist two other variables  $(Z_1 \text{ and } Z_2)$  and two contexts  $(S_1 \text{ and } S_2)$  such that:

<sup>&</sup>lt;sup>9</sup> Definition 2.7.1 was formulated in Pearl (1990) as a relation between events (rather than variables) with the added condition P(Y|X) > P(Y) (in the spirit of Reichenbach 1956, Good 1961, and Suppes 1970). This refinement is applicable to any of the definitions in this section, but it will not be formulated explicitly.