

Recall *algebraic* plausibility measures:

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  - ▶  $\text{Pl}(U \cup V) = \text{Pl}(U) \oplus \text{Pl}(V)$  if  $U \cap V = \emptyset$
  - ▶  $\text{Pl}(U_1 \cap U_2 \mid U_3) = \text{Pl}(U_1 \mid U_2 \cap U_3) \otimes \text{Pl}(U_2 \mid U_3)$  if  $U_2 \cap U_3 \in \mathcal{F}'$ ,  $U_1, U_2, U_3 \in \mathcal{F}$ .

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**Definition:** If  $\text{Pl}$  is algebraic, then  $U$  and  $V$  *do not interact given*  $V'$  if  $\text{Pl}(U \cap V \mid V') = \text{Pl}(U \mid V') \otimes \text{Pl}(V \mid V')$  (if  $V' \in \mathcal{F}'$ ).

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**Lemma:** If  $(W, \mathcal{F}, \mathcal{F}', \text{Pl})$  is an algebraic cps and either  $U \cap V' \in \mathcal{F}'$  or  $V \cap V' \in \mathcal{F}'$ , then  $I_{\text{Pl}}(U, V \mid V')$  implies  $U$  and  $V$  do not interact.

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The converse is not necessarily true.

- ▶ There is an example in the book using possibility measures.
- ▶ We can add an extra condition to get the converse

**Bottom line:** we can separate out the two notions of independence using algebraic plausibility measures.

# Properties of independence for RVs

Recall:

CIRV1[ $\mu$ ]. If  $I_\mu^{rv}(\mathbf{X}, \mathbf{Y} \mid \mathbf{Z})$ , then  $I_\mu^{rv}(\mathbf{Y}, \mathbf{X} \mid \mathbf{Z})$ .

CIRV2[ $\mu$ ]. If  $I_\mu^{rv}(\mathbf{X}, \mathbf{Y} \cup \mathbf{Y}' \mid \mathbf{Z})$ , then  $I_\mu^{rv}(\mathbf{X}, \mathbf{Y} \mid \mathbf{Z})$ .

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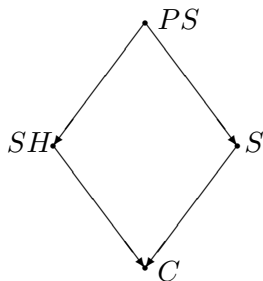
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More general theorem:

**Theorem:** If  $\text{Pl}$  is an algebraic plausibility measure, then these properties continue to hold if we replace  $I_\mu^{rv}$  with  $I_{\text{Pl}}^{rv}$ .

# Qualitative Bayesian Networks

Recall: A *directed acyclic network* consists of a set of nodes and directed edges, where there are no cycles.



- ▶ In a Bayesian network (BN), the nodes are labeled by random variables
- ▶ We can think of the edges as representing causal influence



## More definitions:

- ▶ The *ancestors* of  $X$  in the graph are those random variables that have a potential influence on  $X$ .
  - ▶  $Y$  is an ancestor of  $X$  in graph  $G$  if there is a directed path from  $Y$  to  $X$  in  $G$ —i.e., a sequence  $(Y_1, \dots, Y_k)$  of nodes—such that  $Y_1 = Y$ ,  $Y_k = X$ , and there is a directed edge from  $Y_i$  to  $Y_{i+1}$  for  $i = 1, \dots, k - 1$ .
- ▶ The *parents* of  $X$  in  $G$  ( $\text{Par}_G(X)$ ) are those ancestors of  $X$  directly connected to  $X$ .
  - ▶  $SH$  and  $S$  are the parents of  $C$ ,  $PS$  is the parent of  $S$
- ▶ The *nondescendants* of  $X$  ( $\text{NonDes}_G(X)$ ) are those nodes  $Y$  such that  $X$  is not the ancestor of  $Y$ .

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**Key definition:** The Bayesian network  $G$  (*qualitatively*) represents the probability measure  $\mu$  if, for all nodes  $X$  in  $G$ ,

$$I_{\mu}^{rv}(X, \text{NonDes}_G(X) \mid \text{Par}(X)).$$

- ▶  $X$  is independent of its nondescendants given its parents

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Suppose that a world is characterized by the value of the rvs  $X_1, \dots, X_n$ , and we want to compute the probability of the world  $(x_1, \dots, x_n)$  *without needing to store too many numbers*.

- ▶ Knowing these conditional independencies let's us do this

## An apparent digression: the chain rule

Given arbitrary sets  $U_1, \dots, U_n$ , it is immediate from the definition of conditional probability that

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Applying this observation inductively gives the chain rule:

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Now take  $U_i$  be the event  $X_i = x_i$ .

- the set of all worlds where  $X_i = x_i$

Plugging this into the chain rule gives:

$$\begin{aligned} & \mu(x_1, \dots, x_n) \\ &= \mu(X_1 = x_1 \cap \dots \cap X_n = x_n) \\ &= \mu(X_n = x_n \mid X_1 = x_1 \cap \dots \cap X_{n-1} = x_{n-1}) \times \\ & \quad \mu(X_{n-1} = x_{n-1} \mid X_1 = x_1 \cap \dots \cap X_{n-2} = x_{n-2}) \times \\ & \quad \dots \times \mu(X_2 = x_2 \mid X_1 = x_1) \times \mu(X_1 = x_1). \end{aligned}$$

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To repeat, using the chain rule, we have:

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Now suppose without loss of generality that  $\langle X_1, \dots, X_n \rangle$  is a *topological sort* of (the nodes in)  $G$ .

- ▶ if  $X_i$  is a parent of  $X_j$ , then  $i < j$ .

Thus,  $\{X_1, \dots, X_{k-1}\} \subseteq \text{NonDes}_G(X_k)$ , for  $k = 1, \dots, n$

- ▶ All the descendants of  $X_k$  must have subscripts  $> k$ .
- ▶ Conclusion: all the nodes in  $\{X_1, \dots, X_{k-1}\}$  are independent of  $X_k$  given  $\text{Par}_G(X_k)$ .

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It follows that

$$\begin{aligned} & \mu(X_k = x_k \mid X_{k-1} = x_{k-1} \cap \dots \cap X_1 = x_1) \\ = & \mu(X_k = x_k \mid \cap_{X_i \in \text{Par}(X_k)} X_i = x_i). \end{aligned}$$

So we can greatly simplify our original equation:

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But if  $G$  represents  $\mu$ , then

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**Key point:** If each variable  $X_i$  has relatively few parents, then to compute  $\mu(x_1, \dots, x_n)$ , we need relatively few numbers.

## Quantitative Bayesian Networks

A *quantitative Bayesian network* is a pair  $(G, f)$  consisting of a qualitative Bayesian network  $G$  and a function  $f$  that associates with each node  $X$  in  $G$  a *conditional probability table (cpt)*. If  $\text{Par}_G(X) = \mathbf{Y}$ , then the cpt gives, for each possible setting  $x$  of  $X$  and  $\mathbf{y}$  of  $\mathbf{Y}$ , a number  $f(X, x, \mathbf{Y}, \mathbf{y})$ .

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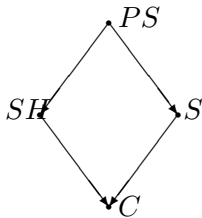
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If  $(G, f)$  quantitatively represents  $\mu$  then we can completely reconstruct  $\mu$  from  $(G, f)$ .

- ▶ Suppose that the world is described by  $N$  binary variables.
- ▶ This means that we are putting a probability distribution on  $2^N$  worlds.
- ▶ But if each rv has at most  $n$  parents, then each cpt requires at most  $2^{n+1}$  numbers
- ▶ At most  $N2^{n+1} \ll 2^N$  numbers needed altogether

**Example:** We get a quantitative BN for smoking by considering the qualitative BN:



together with the following cpts:

$S$	$SH$	$C$
1	1	.6
1	0	.4
0	1	.1
0	0	.01

$PS$	$S$
1	.4
0	.2

$PS$	$SH$
1	.8
0	.3

$PS$
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What about the converse? Given a probability distribution  $\mu$ , can we find a quantitative BN that represents it?

- ▶ Yes! There are lots.

### Construction:

- ▶ Given  $\mu$ , let  $Y_1, \dots, Y_n$  be any permutation of the random variables in  $\mathcal{X}$ .
- ▶ For each  $k$ , find a minimal subset of  $\{Y_1, \dots, Y_{k-1}\}$ , call it  $\mathbf{P}_k$ , such that  $I_\mu^{rv}(\{Y_1, \dots, Y_{k-1}\}, Y_k \mid \mathbf{P}_k)$ .
  - ▶ There is a subset with this property, namely,  $\{Y_1, \dots, Y_{k-1}\}$ .
  - ▶ So there must be a minimal one
- ▶ Add edges from each of the nodes in  $\mathbf{P}_k$  to  $Y_k$ .
- ▶ Call the resulting graph  $G$ .

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**Theorem**  $G$  qualitatively represents  $\mu$ .

- ▶ Now just add the “right” cpts

The Bayesian network constructed depends on the ordering of the edges.

- ▶ Different orderings may lead to different Bayesian networks.
  - ▶ The BN for smoking was constructed with the ordering  $PS, S, SH, C$ .
  - ▶ We could construct another one using the ordering  $C, S, PS, SH$ 
    - ▶ It would have  $C$  at the root
- ▶ Experience has shown that we get “better” BNs if we order the variables causally
  - ▶ If  $X$  has a causal influence on  $Y$ , then  $X$  precedes  $Y$  in the order
    - ▶ This was the case with the original smoking network
  - ▶ “Better” typically means
    - ▶ fewer edges
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    - ▶ easier to elicit the cpt from experts

This construction of BNs used only CIRV1-5

- ▶ Conclusion: it works without change for arbitrary algebraic plausibility measures

# Independencies in BNs

If  $G$  represents  $\mu$ , then an rv in  $G$  is independent of its nondescendants conditional on its parents with respect to  $\mu$ .

- ▶ What other independencies hold?
- ▶ There is a criterion that lets us compute this.

## d-separation

$X$  is *d-separated* ( $d = \text{directed}$ ) from a node  $Y$  by a set  $\mathbf{Z}$  of nodes in  $G$ , written  $d\text{-sep}_G(X, Y \mid \mathbf{Z})$ , if for every *undirected path* from  $X$  to  $Y$  there is a node  $Z'$  on the path such that either

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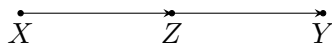
- (b)  $Z' \in \mathbf{Z}$  and has both path arrows leading out; or



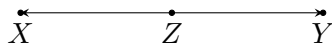
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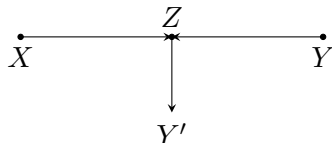
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**Example:**

- ▶  $\{SH, S\}$  d-separates  $PS$  from  $C$ .
- ▶  $\{PS\}$  d-separates  $SH$  from  $S$ .
- ▶  $\{PS, C\}$  does *not* d-separate  $SH$  from  $S$ .

## d-separation: some intuition

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  - ▶  $SH$  and  $S$  are not independent, because they have a common cause ( $PS$ ), but conditioning on  $PS$  makes them independent
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  - ▶  $S$  and  $SH$  are independent conditional on  $PS$ , but they, would become *dependent* if we also conditioned on  $C$

D-separation completely characterizes conditional independence in Bayesian networks:

**Theorem:** If  $\mathbf{X}$  is d-separated from  $\mathbf{Y}$  by  $\mathbf{Z}$  in the Bayesian network  $G$ , then  $I_{\mu}^{rv}(\mathbf{X}, \mathbf{Y} \mid \mathbf{Z})$  holds for all probability measures  $\mu$  compatible with  $G$ . Conversely, if  $\mathbf{X}$  is *not* d-separated from  $\mathbf{Y}$  by  $\mathbf{Z}$ , then there is a probability measure  $\mu$  compatible with  $G$  such that  $I_{\mu}^{rv}(\mathbf{X}, \mathbf{Y} \mid \mathbf{Z})$  does not hold.

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- ▶ The proof of the first half of the theorem requires only CIRV1-5, so holds for all algebraic plausibility measures
- ▶ for the second half, we need some extra conditions.

**Bottom line:** the technology of Bayesian networks can be applied quite widely!