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Knowledge Representation and Inference under Uncertainty

This course requires a basic knowledge of probability theory.

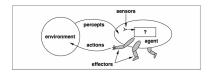
Bayesian neworks are covered by the course textbook, which provides also an informal introduction to probability theory: S. Russell, P. Norvig, *Artificial Intelligence – A Modern Approach*, 4th Ed., Pearson, 2021 (or a previous edition)

For a formal and comprehensive introduction to probability theory, textbooks like the following one are suggested: A.M. Mood, F.A. Graybill, D.C. Boes, *Introduction to the Theory of Statistics*, McGraw-Hill, 1991 / 1998

Introduction

Introduction

Often rational agents must make decisions and act under **uncertainty** about their environments, e.g.:



- the wumpus world (sensors report only local information)
- medical diagnosis from patients' symptoms and outcomes of medical tests
- speech/image recognition from (noisy) audio/video signals
- self-driving vehicles (incomplete and noisy sensory data)

Limitations of logical agents

Logical agents can only deal with propositions that are either true, false, or unknown, but they cannot represent a **degree of belief** about propositions.

Example: consider the agent's knowledge about the configuration of the wumpus world show on the right, after having explored squares (1, 1), (1, 2) and (2, 1).

1,4	2,4	3,4	4,4
1,3	2,3	3,3	4,3
1,2 B OK	2,2		4,2
1,1	2,1 B	3,1	4,1
OK	ок		

A logical agent cannot conclude anything about which of the reachable squares (1,3), (2,2) and (3,1) is **most likely** to be safe from pits: therefore it can only decide the next move **randomly**.

Limitations of logical agents

Diagnosis problems (e.g., in medicine) are a good example of the limitations of the logical approach in dealing with uncertainty.

Example: the following dental diagnosis rule (where p denotes a patient) is wrong, since toothache can have other causes besides cavity:

 $\forall p \; Symptom(p, Toothache) \Rightarrow Disease(p, Cavity)$

Trying to specify **all** possible causes is pointless, e.g.:

 $\forall p \; Symptom(p, Toothache) \Rightarrow$ $Disease(p, Cavity) \lor Disease(p, Abscess) \lor \dots$

Turning the first rule into a **causal** one does not work, either, since not all cavities cause toothache:

 $\forall p \ Disease(p, Cavity) \Rightarrow Symptom(p, Toothache)$

Making rational decisions under uncertainty

Uncertainty can be due to **laziness**, or by **theoretical** or **practical ignorance**.

In domains involving uncertainty (medicine, business, law, etc.), including AI, only a **degree of belief** in sentences of interest can be provided.

Probability theory is a widely used tool to summarise this kind of uncertainty into a numerical value, conventionally in the range [0, 1].

For instance, stating that the probability that a patient with a toothache has a cavity is $0.8\ means$ that

- in all possible situations indistinguishable by the agent from the current one, 80% of patients suffering from toothache have a cavity
- the missing 20% summarises all the other possible causes of toothache the agent is too lazy or ignorant to confirm or deny

Making rational decisions under uncertainty

An effective decision-making agent under uncertainty should have **preferences** about the possible **outcomes** of its actions.

Representing and reasoning with preferences is the subject of **utility theory**.

Rational decision-making under uncertainty can be achieved by **combining** the likelihood (probability) and the preference of actions' outcomes, which in turn is the subject of **decision theory**:

Decision theory = probability theory + utility theory

Basically, a **rational agent** chooses the actions that yields the highest **expected** utility, averaged over all its possible outcomes.

Elements of probability theory

A notation for probability theory

In the following, a version of probability theory suited to AI applications will be presented, as a **formal language** for knowledge representation and reasoning under uncertainty.

This formal language is characterised by two elements:

- the kind of sentence to which degrees of beliefs are assigned
- the distiction between prior and conditional probability statements, related to the evidence available to the agent

The considered notation is an extension of propositional logic.

Propositions and random variables

Degrees of belief are assigned to **propositions** (natural language statements that can be either true or false), that refer to the "state of the world" of interest to the agent.

Propositions used in probability theory describe the "state of the world" in terms of a predefined set of **random variables**, each one referring to a "part" of the world.

Each random variable has a **domain**, i.e., the set of **mutually exclusive** values it can take.

Conventionally, random variables are represented by uppercase names (e.g., *Weather*) and their values as lowercase names (e.g., *sunny*).

The domain of random variables

The domain of random variables can be of three kinds:

- **Boolean**: {*true*, *false*}
- discrete: a set of countable values
 - categorical, e.g.: {*sunny*, *rainy*, *cloudy*, *snow*} for a random variable *Weather*
 - numerical, e.g., integer values
 - Boolean (a particular case of categorical values)
- **continuous**: real numbers, e.g., the interval [0, 1]

This course will consider only **discrete** random variables.

It turns out that also for unordered categorical domains it is convenient to impose an (arbitrary) **ordering** on their values: therefore categorical domains will be written as **tuples**, e.g., $\langle true, false \rangle$, instead of sets.

Elementary and complex propositions

Example: two Boolean random variables related to a given person

- Cavity: whether the lower left wisdom tooth has a cavity
- Toothache: whether that person suffers from toothache

Elementary propositions assert that a **single** random variable has a particular value from its domain, e.g.:

Cavity = true

Complex propositions combine elementary ones using standard logical connectives, e.g.:

Cavity = *true* \land *Toothache* = *false*

Each proposition can be assigned a degree of belief.

Atomic events

Atomic event: a complete description of the "state of the world" about which the agent is uncertain.

It is defined by a proposition that assigns values to **all** the random variables used by the agent.

Example: an agent uses only the two Boolean random variables *Cavity* and *Toothache*

- there are four distinct atomic events
- one of them is:

Cavity = true
$$\land$$
 Toothache = false

Properties of atomic events

- Atomic events are mutually esclusive and exhaustive: exactly one of them must be the case
- - the truth of *Cavity* = *true*
 - the falsehood of $Cavity \Rightarrow Toothache$
- Any proposition is logically equivalent to the disjunction of all atomic events that entail its truth, e.g.:

$$Cavity = true \equiv (Cavity = true \land Toothache = true) \lor (Cavity = true \land Toothache = false)$$

The degree of belief p assigned to a given proposition A, in the absence of any other information, is called **unconditional** or **prior probability**.

The usual notation is P(A) = p. Some examples:

$$P(Weather = sunny) = 0.35$$

 $P(Cavity = true \land Toothache = true) = 0.15$

Probability distribution

Prior probability distribution: the assignment of prior probabilities to **all** the values in the domain of a discrete random variable.

Example: a possible probability distribution for *Weather*, having domain (*sunny*, *rain*, *cloudy*, *snow*):

$$P(Weather = sunny) = 0.7$$

 $P(Weather = rain) = 0.2$
 $P(Weather = cloudy) = 0.08$
 $P(Weather = snow) = 0.02$

In vector notation:

$$P(Wheather) = \langle 0.7, 0.2, 0.08, 0.02 \rangle$$

The axioms of probability

The foundations of probability theory are defined as a set of axioms. In particular, **Kolmogorov's axioms**:

- 1. $0 \leq P(\mathcal{A}) \leq 1$, for any proposition \mathcal{A}
- 2. P(true) = 1, P(false) = 0
 - *true* corresponds to the occurrence of **any** atomic event, which is a **certain** event
 - false corresponds to the occurrence of **no** atomic event, which is an **impossible** event
- 3. $P(A \lor B) = P(A) + P(B) P(A \land B)$, for any pair of propositions A and B

Consequences of the axioms of probability

From the axioms of probability many **theorems** can be derived. **Examples**:

- for any event \mathcal{A} : $P(\mathcal{A}) + P(\neg \mathcal{A}) = 1$
- the values of the probability distribution of a discrete random variable X with domain (x₁, x₂, ..., x_n) must sum to 1 (this is a particular case of the previous theorem):

$$\sum_{i=1}^n P(X=x_i)=1$$

► the probability of a proposition A equals the sum of the probabilities of all the atomic events E_k in which it holds, i.e., if A = V_k E_k, then:

$$P(\mathcal{A}) = \sum_{k} P(\mathcal{E}_{k})$$

Joint probability distribution

It is often useful to consider the probability of the **conjunction** of values of several random variables, e.g.:

$$P(Cavity = true \land Weather = sunny)$$

The probabilities of **all** the combinations of values of a set of random variables is called **joint probability distribution**.

Example: the joint probability distribution P(Cavity, Weather) can be represented by a 4×2 table of probabilities:

		Weather			
		sunny	rain	cloudy	snow
Cavity	true	0.15	0.10	0.03	0.01
	false	0.55	0.10	0.05	0.01

Note that, according to the axioms of probability, the probabilities of a joint distribution sum to 1.

Full joint probability distribution

The joint probability distribution of **all** the random variables used by an agent is called **full joint probability distribution**.

Example: if an agent uses only the variables *Cavity*, *Toothache* and *Weather*, the full joint distribution is

P(Cavity, Toothache, Weather),

which can be represented by a $2 \times 2 \times 4$ table of 16 probabilities. Note that the full joint distribution specifies the probability of

every atomic event.

Conditional probability

If **evidence** about some random variables has been obtained, **conditional** or **posterior** probabilities of the remaining random variables have to be considered.

Example: a patient is **known** to suffer from toothache, and **no other** information is yet available

- the prior probability P(Cavity = true) does not reflect anymore the current state of knowledge
- the conditional probability of the patient's having a cavity given that she suffers from toothache should be considered, e.g., in the conventional notation:

Conditional probability

Conditional probabilities are **defined** in terms of unconditional ones. For any pair of propositions A and B:

$${\sf P}({\cal A}|{\cal B}) riangleq rac{{\sf P}({\cal A} \wedge {\cal B})}{{\sf P}({\cal B})}$$

As a particular case, for any pair of random variables X and Y and any pair of values x and y in the respective domains:

$$P(X = x | Y = y) = \frac{P(X = x, Y = y)}{P(Y = y)}$$

Example:

$$P(Cavity = true | Toothache = true) = \frac{P(Cavity = true, Toothache = true)}{P(Toothache = true)}$$

Conditional probability distribution

For any pair of random variables X and Y and values y in the domain of Y, the **conditional probability distribution** of X given Y = y can be written in vector notation:

$$\mathsf{P}(X|Y=y)$$

Example: $P(Cavity | Toothache = true) = \langle 0.8, 0.2 \rangle$

Note that:

also the values of a conditional distribution sum to 1

every value of the conditioning variable corresponds to a distinct conditional distribution, e.g.:

-
$$P(Cavity|Toothache = true) = \langle 0.8, 0.2 \rangle$$

- $P(Cavity | Toothache = false) = \langle 0.05, 0.95 \rangle$

Product rule

From the **definition** of conditional probability the so called **product rule** immediately follows:

 $P(\mathcal{A} \wedge \mathcal{B}) = P(\mathcal{A}|\mathcal{B})P(\mathcal{B})$

As a particular case, for any pair of random variables X and Y and any pair of values x and y in the respective domains:

$$P(X = x, Y = y) = P(X = x | Y = y)P(Y = y)$$

Their joint distribution can now be rewritten in vector notation:

$$\mathbf{P}(X,Y) = \mathbf{P}(X|Y)\mathbf{P}(Y) ,$$

where the right-hand side denotes element-wise multiplication of the corresponding table values, **not** matrix multiplication.

Marginal probability distribution

Given the joint distribution of any set of random variables $P(X_1, ..., X_n)$, the prior distribution of any subset of them, e.g., $P(X_1, ..., X_m)$, with $1 \le m < n$, is called marginal probability distribution.

Example: Consider the joint distribution **P**(*Cavity*, *Weather*) shown before:

		Weather			
		sunny	rain	cloudy	snow
Cavity	true	0.15	0.10	0.03	0.01
	false	0.55	0.10	0.05	0.01

P(Cavity) and P(Weather) are marginal distributions with respect to P(Cavity, Weather).

Marginal probability distribution

How can one compute a marginal distribution from a **known** joint distribution, e.g., P(Cavity) from P(Cavity, Weather)?

To this aim, a result mentioned above can be exploited, i.e.: the probability of any proposition equals the sum of the probabilities of all the atomic events in which it holds.

For instance, P(Cavity = true) is the sum of all the values of P(Cavity, Weather) where Cavity = true:

P(Cavity = true) = 0.15 + 0.10 + 0.03 + 0.01 = 0.39

Marginalisation, or sum rule

In general, for any **disjoint** sets of variables **Y** and **Z**:

$$\mathsf{P}(\mathsf{Y}) = \sum_{\mathsf{z}} \mathsf{P}(\mathsf{Y},\mathsf{Z}=\mathsf{z}) \;,$$

where the sum is over all combinations of values of the variables Z.

This process is called marginalisation, or sum rule, and can be applied if the joint distribution P(Y, Z) is known.

It is often useful to rewrite the sum rule in terms of the **product** rule, using **conditional** probabilities:

$$\mathbf{P}(\mathbf{Y}) = \sum_{\mathbf{z}} \mathbf{P}(\mathbf{Y}|\mathbf{Z} = \mathbf{z}) P(\mathbf{Z} = \mathbf{z})$$

Probabilistic inference

In most cases of interest for rational agents in AI, **probabilistic inference** consists in computing the **posterior** probability of a set of **query** variables **Q** given the **evidence** about a distinct set of variables $\mathbf{E} = \mathbf{e}$

- either for a single value of interest \mathbf{q} : $P(\mathbf{Q} = \mathbf{q} | \mathbf{E} = \mathbf{e})$
- or the posterior distribution P(Q|E = e)

If the **full** joint distribution of the variables used by an agent is **known**, the **sum** and **product** rules allow the agent, in priniciple, to carry out **any** probabilistic inference, as shown in the following.

Inference using the full joint distribution: example

An agent uses three Boolean variables related to a given patient of a dentist:

- Toothache and Cavity, defined previously
- Catch: the dentist's steel probe catches in the lower left wisdom tooth

Assume the full joint distribution is:

	Tootha	che = t	Toothache = f		
	Catch = t	Catch = f	Catch = t	Catch = f	
Cavity = t	0.108	0.012	0.072	0.008	
Cavity = f	0.016	0.064	0.144	0.576	

Inference using the full joint distribution: example

Unconditional probabilities P(Q), such as

$$P(Cavity = true \lor Toothache = true)$$
,

can be computed as shown above, as the sum of the probabilities of the atomic events in which the event ${\cal Q}$ holds:

$$P(Cavity = true \lor Toothache = true) \\= 0.108 + 0.012 + 0.072 + 0.008 + 0.016 + 0.064 = 0.28$$

Inference using the full joint distribution: example

Conditional probabilities $P(Q|\mathcal{E})$, such as

can be computed using the definition of conditional probability, in terms of **unconditional** ones, which can in turn be computed from the full joint distribution as shown above:

$$P(Cavity = t | Toothache = t) = \frac{P(Cavity = t \land Toothache = t)}{P(Toothache = t)}$$
$$= \frac{0.108 + 0.012}{0.108 + 0.012 + 0.016 + 0.064}$$

Note that the last step corresponds to applying the **sum rule** to the **full** joint distribution.

Inference using the full joint distribution

A general inference procedure for conditional distributions P(Q|E = e), for the case of a single query variable Q, with Y denoting all the remaining variables:

- 1. rewriting P(Q|E = e) using the definition of conditional probability, in terms of **unconditional** probabilities
- 2. using the sum rule to compute the numerator and denominator, from the full joint distribution P(Q, E, Y)

$$\mathsf{P}(Q|\mathsf{E}=\mathsf{e}) = \frac{\mathsf{P}(Q,\mathsf{E}=\mathsf{e})}{\mathsf{P}(\mathsf{E}=\mathsf{e})} = \frac{\sum_{\mathsf{y}} \mathsf{P}(Q,\mathsf{E}=\mathsf{e},\mathsf{Y}=\mathsf{y})}{\sum_{q,\mathsf{y}} \mathsf{P}(Q=q,\mathsf{E}=\mathsf{e},\mathsf{Y}=\mathsf{y})}$$

Inference using the full joint distribution

To sum up, if the **full joint** distribution of the random variables of interest is known, the probability of **any** event can be computed using

- the definition of conditional probability
- the sum and product rules

In practice, two issues arise:

- **how** to compute or estimate the full joint distribution?
- what is the computational complexity of the inference procedure based on it?

Defining the full joint distribution

Defining the full joint distribution of variables X_1, \ldots, X_n amounts to **assigning** a probability value to **each** of the **atomic events**

$$X_1 = x_1 \wedge \ldots \wedge X_n = x_n$$

To assign a probability value to **any** event, including atomic ones, three different approaches exist, based on different **definitions** of probability:

- classical
- frequentist
- subjective

Classical probability

In some cases the **atomic** events that describe the "state of the world" can be assumed to be:

- mutually exclusive
- equally likely
- random

Examples: tossing a coin, throwing a dice.

In such cases the **classical** (or a **priori**) definition of probability can be applied for **any** event A:

 $P(\mathcal{A}) = \frac{\text{number of atomic events where } \mathcal{A} \text{ holds}}{\text{total number of atomic events}}$

Classical probability: examples

Tossing a coin: probability of a head (or a tail)?

- Throwing a dice
 - probability of face 3 up?
 - probability of an even face up?
- Tossing two coins: probability of getting two heads?
- Throwing two dice
 - probability of getting (6,6)?
 - probability that the sum of the faces up is 6?
- Picking a card from a well shuffled deck of 52 cards
 - probability of picking an ace of hearts (\heartsuit)?
 - probability of picking an ace or a spade (\spadesuit) ?

Frequentist probability

In most practical applications the assumptions of classical probability are **not valid**, e.g., computing P(A) for:

- A = face 6 up after throwing a **loaded** dice
- A = a chip manufactured by company XYZ is faulty
- A = a MSc student in CECAI graduates within 2 years

Nevertheless, it may be possible to observe **multiple** "states of the world" under **similar** and **uniform** conditions, e.g.:

- repeatedly throwing a loaded dice and observing the outcomes
- testing a sample of the chips manufactured by company XYZ
- collecting records of past MSc students in CECAI

Frequentist probability

If **multiple** "states of the world" can be observed under **similar** and **uniform** conditions, the **frequentist** (or **a posteriori**) probability definition can be used.

It estimates the probability of any event A, including atomic events, as the relative frequency of its occurrences:

 $P(\mathcal{A}) = \frac{\text{number of observations where } \mathcal{A} \text{ occurs}}{\text{total number of observations}}$

Examples

- fraction of throws of a dice leading to face 6 up
- fraction of XYZ's sampled chips that are faulty
- fraction of past MSc students in CECAI who graduated within 2 years

Subjective probability

Common requirement of classical and frequentist probability: a **conceptual** "experiment" in which the outcomes can occur under **uniform conditions**.

This requirement is not always fulfilled. For instance, how to compute the probability that

- the piano player John Smith will break one or both of his hands within the next 10 years?
- the third World War will start within 2025?

In such cases **subjective** probability can be used, e.g., involving domain experts' judgement.

Computational complexity of probabilistic inference

For *n* discrete random variables X_1, \ldots, X_n whose domains are made up of d_1, \ldots, d_n values, the number of atomic events is

$$d_1 \times \ldots \times d_n$$

Therefore, the number of probability values to be assigned to define the full joint distribution (they must sum to 1) grows **exponentially** in the domain size:

$$d_1 imes \ldots imes d_n - 1$$

Example: for *n* Boolean variables

- the number of atomic events is 2ⁿ
- the number of probability values to assign is $2^n 1$

Computational complexity of probabilistic inference

Also the probabilistic inference procedure presented above has an **exponential** complexity in the domain size, due to the **sum rule**.

For instance, for *n* Boolean variables X_1, \ldots, X_n the marginal distribution $\mathbf{P}(X_1)$ is given by

$$\sum_{x_2,\ldots,x_n \in \{0,1\}^{n-1}} \mathbf{P}(X_1, X_2 = x_2, \ldots, X_n = x_n) ,$$

which requires n - 2 sums for **each** of the 2^{n-1} combinations of values x_2, \ldots, x_n , i.e., $\mathcal{O}(2^n)$ sums.

Therefore, the full joint distribution in tabular form is not a practical tool for building probabilistic reasoning systems.

Independence

Two events A and B are said to be **independent**, if **any** of the following **equivalent** relation holds:

$$P(\mathcal{A}|\mathcal{B}) = P(\mathcal{A}), \quad P(\mathcal{B}|\mathcal{A}) = P(\mathcal{B}), \quad P(\mathcal{B} \land \mathcal{A}) = P(\mathcal{A})P(\mathcal{B})$$

In particular, two variables X and Y are independent, if any of the following relation holds between their **distributions**:

$$\mathbf{P}(X|Y) = \mathbf{P}(X), \quad \mathbf{P}(Y|X) = \mathbf{P}(Y), \quad \mathbf{P}(X,Y) = \mathbf{P}(X)\mathbf{P}(Y)$$

Independence is useful to our purposes since it reduces

- the number of probability values to be assigned to specify the full joint distribution
- the computational complexity of probabilistic inference

Independence: example

Assume that an agent uses the variables

- Toothache, Catch and Cavity (all Boolean)
- Weather, with domain (sunny, rain, cloudy, snow)

The full joint distribution P(Toothache, Catch, Cavity, Weather) is specified by a $2 \times 2 \times 2 \times 4$ table of 32 probability values.

Is there any way to **reduce** the number of probabilty values to assign, to specify this distribution?

To this aim, one may wonder whether and how dental problems (toothache, the dentist's probe catching in a tooth, and cavities) and the weather are related to each other...

Independence: example

... Consider rewriting the full joint distribution using the product rule:

P(*Toothache*, *Catch*, *Cavity*, *Weather*)

= **P**(*Weather*|*Toothache*, *Catch*, *Cavity*)**P**(*Toothache*, *Catch*, *Cavity*)

It is reasonable to assume that dental problems do not influence the weather, i.e., the variable *Weather* can be assumed to be **independent** from *Toothache*, *Catch* and *Cavity*:

 $\mathbf{P}(Weather | Toothache, Catch, Cavity) = \mathbf{P}(Weather)$

This allows rewriting the full joint distribution as:

P(*Toothache*, *Catch*, *Cavity*, *Weather*)

 $= \mathbf{P}(Weather)\mathbf{P}(Toothache, Catch, Cavity)$

Independence: example

Using the last expression, the full joint distribution can be specified by:

- $P(Weather): 4^1 1 = 3$ probability values,
- ▶ **P**(*Toothache*, *Catch*, *Cavity*): $2^3 1 = 7$ probability values,

for a total of 3 + 7 = 10 values, instead of $2^4 - 1 = 31!$

In the **best case**, **all** variables are independent. For instance, the full joint distribution of n Boolean variables is specified by

- ▶ $2^n 1 = O(2^n)$ values, in general (exponential in *n*)
- ▶ n × (2¹ − 1) = n = O(n) values, if they are independent (linear in n), i.e.:

$$\mathbf{P}(X_1,\ldots,X_n)=\prod_{i=1}^n\mathbf{P}(X_i)$$

Unfortunately, in practice **absolute** independence is quite rare, even among subsets of random variables.

Consider the joint distribution P(Toothache, Catch, Cavity), which is specified by $2^3 - 1 = 7$ values.

These variables, or subsets of them, **cannot** be assumed to be independent. For instance, if the probe catches in the tooth, probably there is a cavity, and that probably causes a toothache.

Noting that cavity is a possible **cause** of **effects** like toothache and the dentist's probe catching in a tooth, let us rewrite the joint distribution using the **product rule** in the form P(effect|cause):

P(Toothache, Catch, Cavity) = P(Toothache, Catch|Cavity)P(Cavity)

What can we say about **P**(*Toothache*, *Catch*|*Cavity*)?

Given the state (presence of absence) of a cavity, *Toothache* and *Catch* have no direct influence on each other, and therefore can be assumed to be **independent**, which is written as:

P(Toothache, Catch|Cavity) = P(Toothache|Cavity)P(Catch|Cavity)Equivalently:

P(*Toothache*|*Catch*, *Cavity*) = **P**(*Toothache*|*Cavity*)

This is an example of conditional independence.

This allows the joint distribution to be rewritten as

$$\begin{split} \mathsf{P}(\textit{Toothache},\textit{Catch},\textit{Cavity}) &= \mathsf{P}(\textit{Toothache},\textit{Catch}|\textit{Cavity})\mathsf{P}(\textit{Cavity}) \\ &= \mathsf{P}(\textit{Toothache}|\textit{Cavity})\mathsf{P}(\textit{Catch}|\textit{Cavity})\mathsf{P}(\textit{Cavity}) \end{split}$$

How many probability values must be assigned to specify the joint distribution P(Toothache, Catch, Cavity), using the conditional independence assumption above?

- **P**(*Toothache*|*Cavity*): $2 \times (2^1 1) = 2$ values,
- $P(Catch|Cavity): 2 \times (2^1 1) = 2$ values,
- ► P(Cavity): 2¹ − 1 = 1 value,

for a total of 2 + 2 + 1 = 5 values, instead of 7.

This may seem a small gain...

... However, in practical applications involving many random variables and non-Boolean domains the advantage can be notable.

For instance, consider *n* Boolean variables X_1, \ldots, X_n conditionally independent given a Boolean variable *Y*:

- ► the full joint distribution P(X₁,...,X_n,Y) is specified by 2ⁿ⁺¹ - 1 = O(2ⁿ⁺¹) probability values
- the equivalent expression $\mathbf{P}(Y)\mathbf{P}(X_1|Y) \times \ldots \times \mathbf{P}(X_n|Y)$ is specified by

-
$$P(Y)$$
: $2^1 - 1 = 1$ value,
- $P(X_i|Y), i = 1, ..., n$: $n \times (2^1 - 1) = n$ values,

for a total of $n + 1 = \mathcal{O}(\mathbf{n})$ values

This means that the number of probability values to assign reduces from **exponential** to **linear** in the number of variables.

In general, two variables X and Y are **conditionally independent** given a variable Z, if:

$$\mathbf{P}(X, Y|Z) = \mathbf{P}(X|Z)\mathbf{P}(Y|Z)$$

As with absolute independence, this is equivalent to:

$$\mathbf{P}(X|Y,Z) = \mathbf{P}(X|Z)$$
 and $\mathbf{P}(Y|X,Z) = \mathbf{P}(Y|Z)$

Conditional independence is very useful in practice:

- it allows probabilistic systems to scale up
- it is much more common than absolute independence

Probabilistic inference

As mentioned above, probabilistic inference in AI usually consists in computing **posterior** probabilities $P(Q|\mathcal{E})$

- Q: a query event
- \blacktriangleright \mathcal{E} : an available evidence

In terms of random variables, the most general form of probabilistic inference consists in computing a **conditional** probability distribution P(Q|E)

- Q: a set of query variables
- **E**: a distinct set of **evidence** variables

Probabilistic inference: Bayes' rule

Consider any pair of events A and B. The probability of their conjunction can be rewritten using the product rule:

$$P(\mathcal{A} \wedge \mathcal{B}) = P(\mathcal{A}|\mathcal{B})P(\mathcal{B})$$

 $P(\mathcal{B} \wedge \mathcal{A}) = P(\mathcal{B}|\mathcal{A})P(\mathcal{A})$

Since $P(A \land B) = P(B \land A)$, it follows that:

$$P(\mathcal{B}|\mathcal{A}) = rac{P(\mathcal{A}|\mathcal{B})P(\mathcal{B})}{P(\mathcal{A})}$$

The above equation is known as **Bayes' rule**. It turns out to be a fundamental tool for probabilistic inference in Al systems.

Probabilistic inference: Bayes' rule

In terms of probability distributions of random variables, Bayes' rule can be written as:

$$\mathsf{P}(X|Y) = \frac{\mathsf{P}(Y|X)\mathsf{P}(X)}{\mathsf{P}(Y)}$$

The most general form involves sets of random variables:

$$\mathbf{P}(\mathbf{X}|\mathbf{Y}) = \frac{\mathbf{P}(\mathbf{Y}|\mathbf{X})\mathbf{P}(\mathbf{X})}{\mathbf{P}(\mathbf{Y})}$$

Note that the denominator can be rewritten in turn as a function of the same distributions in the numerator, using the sum and product rules:

$$\mathbf{P}(\mathbf{Y}) = \sum_{\mathbf{x}} \mathbf{P}(\mathbf{Y}, \mathbf{X} = \mathbf{x}) = \sum_{\mathbf{x}} \mathbf{P}(\mathbf{Y} | \mathbf{X} = \mathbf{x}) P(\mathbf{X} = \mathbf{x})$$

When and why is Bayes' rule useful?

To compute **one** conditional probability, Bayes' rule requires **three** other probabilities. So, why is it useful in AI (and in other domains)?

The reason is that probabilistic inference is usually of **diagnostic** form:

P(cause|effect),

where the evidence consists of observed **effects** (e.g., a symptom like toothache) of a possible **cause** of interest (e.g., a cavity).

In practice, **diagnostic knowledge**, i.e., estimating P(cause|effect), is more difficult to obtain than **causal knowledge**, i.e., P(effect|cause).

For instance, P(Cavity|Toothache) can be more difficult to estimate than P(Toothache|Cavity).

(cont.)

When and why is Bayes' rule useful?

In this context, Bayes' rule allows to carry out diagnostic inference using **only** causal knowledge:

$$P(cause|effect) = rac{P(effect|cause)P(cause)}{P(effect)}$$

Note that, as shown above, P(effect) can be computed in terms of P(effect|cause) and P(cause): therefore, only these two probabilities need to be estimated.

Probabilistic inference using Bayes' rule: an example

A doctor knows that

meningitis causes stiff neck in 50% of patients (causal knowledge)

- 1 out of 50.000 individuals in the population has meningitis
- 1 out of 20 individuals suffers from stiff neck

Using the Boolean variables M and S to denote whether a random individual has meningitis and stiff neck, respectively, the above knowledge can be formalised as:

$$\blacktriangleright P(S = true | M = true) = 0.5$$

•
$$P(M = true) = 1/50.000$$

•
$$P(S = true) = 1/20$$

(cont.)

Probabilistic inference using Bayes' rule: an example

The doctor can be interested in computing the probability that a patient with stiff neck has meningitis (**diagnostic inference**).

She can exploit the above knowledge, using Bayes' rule:

$$P(M = true|S = true) = \frac{P(S = true|M = true)P(M = true)}{P(S = true)}$$
$$= \frac{0.5 \times 1/50.000}{1/20} = 0.0002$$

This is a particular case of the general fact mentioned above: in the medical domain, **diagnostic knowledge** (from symptoms to diseases) is usually more difficult to obtain than **causal knowledge** (from diseases to symptoms).

(cont.)

Probabilistic inference using Bayes' rule: an example

Even if diagnostic knowledge is available from statistical observation to directly estimate P(M = true|S = true), what happens if there is an **epidemic** of meningitis, i.e., if P(M = true) increases?

Intuitively, also P(M = true | S = true) will increase, but how to update it?

Also in this case Bayes' rule is useful, because:

- P(S = true | M = true) is **unaffected** by the epidemic
- thus P(M = true | S = true) increases **proportionately** with P(M = true)

This is an example of how **causal knowledge** provides the necessary **robustness** to make probabilistic systems feasible in practical applications.

Consider again the variables Toothache, Catch and Cavity.

A dentist may collect **two** evidences: her probe chatches in the patient's aching tooth. She is therefore interested in the following conditional distribution:

P(*Cavity* | *Toothache*, *Catch*)

This is a kind of diagnostic inference of the form

P(cause|effects)

We already know that using the full joint distribution (if known) to compute the above conditional probability does not scale up to large numbers of variables.

The doctor can try to apply Bayes' rule to turn diagnostic inference into **causal** inference of the form P(effects|cause):

$$\begin{split} \textbf{P}(\textit{Cavity}|\textit{Toothache},\textit{Catch}) \\ &= \frac{\textbf{P}(\textit{Toothache},\textit{Catch}|\textit{Cavity})\textbf{P}(\textit{Cavity})}{\textbf{P}(\textit{Toothache},\textit{Catch})} \end{split}$$

However, estimating P(Toothache, Catch|Cavity) still requires $2 \times (2^2 - 1) = 6$ probability values, which does not scale up to larger numbers of evidence and query variables, either.

For instance, for *n* evidence variables E_1, \ldots, E_n and a single query variable Q, all Boolean, the conditional distributions $\mathbf{P}(E_1, \ldots, E_n | Q)$ require $2 \times (2^n - 1)$ probability values.

(cont.)

Fortunately, we have seen that a **conditional independence** assumption can be made:

P(Toothache, Catch|Cavity) = P(Toothache|Cavity)P(Catch|Cavity)

Each distribution in the right-hand side is specified by

$$2\times(2^1-1)=2$$

probability values, for a total of 4 values instead of 6.

In general, if *n* Boolean evidence variables E_1, \ldots, E_n are conditionally independent on a Boolean query variable Q, i.e.:

$$\mathbf{P}(E_1,\ldots,E_n|Q)=\prod_{i=1}^n\mathbf{P}(E_i|Q)\;,$$

the number of probability values to specify reduces from $2 \times (2^n - 1) = \mathcal{O}(2^{n+1})$ to $n \times 2 \times (2^1 - 1) = 2n = \mathcal{O}(n)$.

This example highlights the usefulness of **conditional independence**, together with Bayes' rule, for **diagnostic** inference of the form P(cause|effects) involving the combination of different pieces of evidence.

Exercise: the wumpus world revisited

Two properties of the wumpus world are that

- a pit causes breezes in all neighboring squares
- each square other than (1,1) can contain a pit with probability 0.2

Consider now the agent's knowledge about the configuration of the wumpus world shown on the right, after having explored squares (1, 1), (1, 2) and (2, 1).

1,4	2,4	3,4	4,4
1,3	2,3	3,3	4,3
1,2 В ОК	2,2		4,2
1,1	2,1 B	3,1	4,1
OK	OK		

Based on the current agent's knowledge, each of the three reachable squares (1,3), (2,2) and (3,1) can contain a pit: what is the probability that each of them contains a pit?

Bayesian networks

Probabilistic graphical models

Probabilistic graphical models: a framework to represent in graphical form the **structure** of probability distributions.

Main classes of graphical models:

- Bayesian networks: directed acyclic graphs
- Markov random fields: undirected graphs

This course focuses on Bayesian networks, which are useful for

- representing the joint distribution of random variables, expressing causal dependencies and conditional independence relations between them
- developing efficient probabilistic inference algorithms based on graph structure

The chain rule

Bayesian networks exploit a particular expression of the joint distribution of a set of variables based on the **product rule**.

Example: consider three variables X_1, X_2, X_3 ; their joint distribution $P(X_1, X_2, X_3)$ can be rewritten using the product rule as:

 $\mathbf{P}(X_1, X_2, X_3) = \mathbf{P}(X_3 | X_2, X_1) \mathbf{P}(X_2, X_1)$

Applying again the product rule to $P(X_2, X_1)$:

$$\mathbf{P}(X_1, X_2, X_3) = \mathbf{P}(X_3 | X_2, X_1) \mathbf{P}(X_2 | X_1) \mathbf{P}(X_1)$$

Different but **equivalent** expressions of the **same** joint distribution can be obtained considering a different **order** between the variables, e.g.:

$$\mathbf{P}(X_1, X_2, X_3) = \mathbf{P}(X_2 | X_1, X_3) \mathbf{P}(X_1 | X_3) \mathbf{P}(X_3)$$

The chain rule

In general, for a given order of n variables:

$$P(X_1, X_2, ..., X_n) = P(X_n | X_{n-1}, ..., X_1) P(X_{n-1} | X_{n-2}, ..., X_1) \cdots P(X_2 | X_1) P(X_1)$$

= $\prod_{k=1}^{n} P(X_k | X_{k-1}, ..., X_1)$

This equivalence is called chain rule.

Note the **structure** of this expression of the joint distribution:

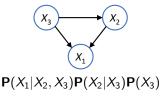
- a product of n 1 conditional and 1 unconditional distributions
- each term is the distribution of a single, distinct variable
- each conditional distribution is conditioned on all the variables that follow the considered one in the chosen order

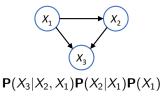
Bayesian network structure

The expression of the joint distribution obtained using the **chain rule** can be represented by a **Bayesian network** (BN), which is a **directed acyclic graph** (DAG) where

- each node represents one of the random variables, and is associated with its distribution in the expression of the chain rule
- oriented edges represent conditional dependencies, linking each variable (node) with the ones on which its distribution is conditioned

Example: BNs representing two possible, **equivalent** expressions of the joint distribution $P(X_1, X_2, X_3)$



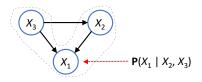


Bayesian network structure

In a BN the conditional distribution $\mathbf{P}(X_k|X_{k-1},\ldots,X_1)$ is represented by

- \blacktriangleright a **node** associated with X_k
- edges pointing from nodes X_{k-1},...,X₁ to X_k, which are called the parents of X_k





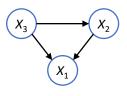
Denoting the set of parents of X_k by $pa(X_k)$, the chain rule can be concisely written as

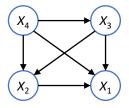
$$\mathsf{P}(X_1,X_2,\ldots,X_n) = \prod_{k=1}^n \mathsf{P}(X_k|\mathit{pa}(X_k))$$

Bayesian network structure

A BN representing a joint distribution expressed through the chain rule is a **fully connected** DAG, i.e., there is an edge (regardless of the direction) between **every** pair of nodes.

Examples:





 $\begin{aligned} \mathbf{P}(X_1, X_2, X_3, X_4) &= \\ \mathbf{P}(X_1 | X_2, X_3, X_4) \mathbf{P}(X_2 | X_3, X_4) \times \\ \mathbf{P}(X_3 | X_4) \mathbf{P}(X_4) \end{aligned}$

Probabilistic inference with Bayesian networks

The chain rule suggests that, instead of estimating the full joint distribution, one could estimate the corresponding conditional (or marginal) distributions, and use them for probabilistic inference.

Example: given three variables X_1, X_2, X_3 , we know that any probability involving them can be computed from the full joint distribution using the definition of conditional probability and the sum rule, e.g.:

$$\mathbf{P}(X_1|X_2) = \frac{\mathbf{P}(X_1, X_2)}{\mathbf{P}(X_2)} = \frac{\sum_{x_3} \mathbf{P}(X_1, X_2, X_3 = x_3)}{\sum_{x_1, x_3} \mathbf{P}(X_1 = x_1, X_2, X_3 = x_3)}$$

Using the chain rule, the numerator could be computed, e.g., as:

$$\sum_{x_3} \mathbf{P}(X_1|X_2, X_3 = x_3) \mathbf{P}(X_2|X_3 = x_3) \mathbf{P}(X_3 = x_3)$$

Complexity of probabilistic inference with BNs

Does the chain rule provide any advantage over the full joint distribution?

If one considers the number of probability values to estimate for specifying the distributions required by the chain rule, the answer is **no**: both approaches are **equivalent** under this aspect.

Example: consider three Boolean variables, and one possible expression of the chain rule:

$$\mathbf{P}(X_1, X_2, X_3) = \mathbf{P}(X_1 | X_2, X_3) \mathbf{P}(X_2 | X_3) \mathbf{P}(X_3)$$

To estimate $\mathbf{P}(X_1, X_2, X_3)$ one needs to specify $2^3 - 1 = 7$ probability values.

(cont.)

Complexity of probabilistic inference with BNs

To estimate the three distributions required by the chain rule, one needs to specify:

•
$$P(X_1|X_2, X_3): 2^2 \times (2^1 - 1) = 4$$
 values,

•
$$P(X_2|X_3): 2 \times (2^1 - 1) = 2$$
 values,

•
$$P(X_3): 2^1 - 1 = 1$$
 value,

for a total of 7 values, as for the full joint distribution.

On the other hand, we have seen that **conditional independence** can reduce the complexity of estimating conditional distributions.

Conditional independence relations can be exploited in the expression of the chain rule. For instance, consider the above example on three Boolean variables:

$$\mathbf{P}(X_1, X_2, X_3) = \mathbf{P}(X_1 | X_2, X_3) \mathbf{P}(X_2 | X_3) \mathbf{P}(X_3)$$

If X_1 is conditionally independent on X_2 given X_3 , i.e.,

$$\mathbf{P}(X_1|X_2,X_3) = \mathbf{P}(X_1|X_3)$$
,

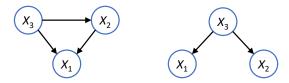
to estimate $P(X_1|X_3)$ one needs to specify $2 \times (2^1 - 1) = 2$ probability values instead of $2^2 \times (2^1 - 1) = 4$.

Conditional independence relations are expressed in the structure of a BN by the **absence** of the corresponding edges, leading to a DAG which is **not** fully connected.

In the previous example, the **general** expression of the chain rule corresponds to the fully connected BN shown below on the left. The expression resulting from the conditional independence assumption is:

$$\mathbf{P}(X_1, X_2, X_3) = \mathbf{P}(X_1 | X_3) \mathbf{P}(X_2 | X_3) \mathbf{P}(X_3) ,$$

which corresponds to the BN below on the right.

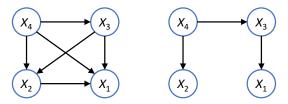


Example: a possible expression of the full joint distribution of four Boolean variables, corresponding to the BN shown below on the left:

 $\mathbf{P}(X_1, X_2, X_3, X_4) = \mathbf{P}(X_1 | X_2, X_3, X_4) \mathbf{P}(X_2 | X_3, X_4) \mathbf{P}(X_3 | X_4) \mathbf{P}(X_4)$

If X_1 is conditionally independent on X_2 and X_4 given X_3 , and X_2 is conditionally independent on X_3 given X_4 (see the BN below on the right):

 $\mathbf{P}(X_1, X_2, X_3, X_4) = \mathbf{P}(X_1 | X_3) \mathbf{P}(X_2 | X_4) \mathbf{P}(X_3 | X_4) \mathbf{P}(X_4)$



(cont.)

How many proability values need to be specified to estimate the full joint distribution using the chain rule?

- General form: $\mathbf{P}(X_1|X_2, X_3, X_4)\mathbf{P}(X_2|X_3, X_4)\mathbf{P}(X_3|X_4)\mathbf{P}(X_4)$
 - $\mathbf{P}(X_1|X_2,X_3,X_4)$: $2^3 \times (2^1 1) = 8$ values,
 - $\mathbf{P}(X_2|X_3,X_4)$: 2² × (2¹ 1) = 4 values,
 - $\mathbf{P}(X_3|X_4)$: 2 × (2¹ 1) = 2 values,
 - $\mathbf{P}(X_4)$: $2^1 1 = 1$ value, for a total of **15** values
- Taking into account conditional independence relations: $\mathbf{P}(X_1|X_3)\mathbf{P}(X_2|X_4)\mathbf{P}(X_3|X_4)\mathbf{P}(X_4)$

-
$${f P}(X_1|X_3)$$
: 2 $imes$ (2¹ – 1) = 2 values,

- $\mathbf{P}(X_2|X_4)$: 2 × (2¹ 1) = 2 values,
- $\mathbf{P}(X_3|X_4)$: 2 × (2¹ 1) = 2 values,
- $\mathbf{P}(X_4)$: $2^1 1 = 1$ value, for a total of **7** values

To efficiently represent full joint distributions through the chain rule and the corresponding BNs, conditional independence relations should be exploited whenever possible.

How can such relations be identified, and exploited?

Main issue: the chain rule provides n! **equivalent** expressions of the full joint distribution of n variables, one for each possible **order** between them. What is the "best" order to choose?

Example: two possible expressions of the chain rule for the full joint distribution of four variables, $P(X_1, X_2, X_3, X_4)$:

 $\mathbf{P}(X_1|X_2, X_3, X_4) \mathbf{P}(X_2|X_3, X_4) \mathbf{P}(X_3|X_4) \mathbf{P}(X_4)$ $\mathbf{P}(X_4|X_3, X_2, X_1) \mathbf{P}(X_3|X_2, X_1) \mathbf{P}(X_2|X_1) \mathbf{P}(X_1)$

If one knows that:

- ► X_1 is conditionally independent on X_2 and X_4 given X_3 : $\mathbf{P}(X_1|X_2, X_3, X_4) = \mathbf{P}(X_1|X_3)$,
- ► X_2 is conditionally independent on X_3 given X_4 : $\mathbf{P}(X_2|X_3, X_4) = \mathbf{P}(X_2|X_4),$

the first expression above can be simplified as:

$$\mathbf{P}(X_1|X_3)\mathbf{P}(X_2|X_4)\mathbf{P}(X_3|X_4)\mathbf{P}(X_4)$$
,

whereas the second one cannot.

To identify the "best" order among variables before applying the chain rule, two facts can be exploited:

- causal knowledge in the form P(effect|causes) is often easier to obtain than diagnostic knowledge, i.e., P(causes|effect)
- conditional independence relations turn out to be more easily identifiable in conditional distributions representing causal knowledge, i.e., P(effects|cause)

Example: we have seen that *Toothache* and *Catch* can be considered conditionally independent given *Cavity*:

P(*Toothache*|*Catch*, *Cavity*) = **P**(*Toothache*|*Cavity*)

If the chain rule is applied to their full joint distribution as:

$$\begin{split} \mathsf{P}(\textit{Toothache},\textit{Catch},\textit{Cavity}) &= \mathsf{P}(\textit{Toothache},\textit{Catch}|\textit{Cavity})\mathsf{P}(\textit{Cavity}) \\ &= \mathsf{P}(\textit{Toothache}|\textit{Catch},\textit{Cavity})\mathsf{P}(\textit{Catch}|\textit{Cavity})\mathsf{P}(\textit{Cavity}) , \end{split}$$

it can be simplified thanks to the above assumption into:

P(*Toothache*|*Cavity*)**P**(*Catch*|*Cavity*)**P**(*Cavity*)

Note that the order of the variables used to apply the chain rule leads to the term in the form P(effects|cause) highlighted in red, which allows exploiting the conditional independence assumption.

(cont.)

BNs corresponding to the first, general expression of the full joint distribution P(Toothache, Catch, Cavity) through the chain rule (left) and to the simplified expression obtained from the conditional independence assumption (right).



The above is an example of the **general rule** to choose the "best" order between the variables when applying the chain rule:

select the "root cause" variables first, then the ones they **directly** influence, and so on, until reaching the variables which have **no direct causal influence** on the others.

The same rule allows to **directly** construct a BN by adding nodes one at a time, together with the corresponding edges, without writing the expression of the chain rule first.

You have a new burglar alarm installed at home.

It is fairly reliable at detecting a burglary, but also responds on occasion to minor earthquakes.

You also have two neighbors, John and Mary, who have promised to call you at work when they hear the alarm.

John always calls when he hears the alarm, but sometimes confuses the telephone ringing with the alarm and calls then, too. Mary, on the other hand, likes rather loud music and sometimes misses the alarm altogether.

You may be interested in different probabilistic inferences, e.g., estimating the probability of a burglary given the evidence of who has or has not called.

taken from Russell and Norvig, Artificial Intelligence - A modern Approach, 2nd Ed., Pearson, 2003

First step: what random variables should be used to represent the events of interest?

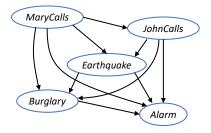
A possible choice is the following set of Boolean variables, related to events occurring over any whole day

- ▶ Alarm (A for brevity): whether the alarm sounded or not
- Burglary (B): whether a burglary occurred or not
- ► *Earthquake* (*E*): whether an earthquake occurred or not
- ► JohnCalls (J): whether John called or not
- MaryCalls (M): whether Mary called or not

One of the possible expressions of the full joint distribution obtained using the chain rule is, e.g.:

 $\mathbf{P}(A, B, E, J, M)$ = $\mathbf{P}(A|B, E, J, M)\mathbf{P}(B|E, J, M)\mathbf{P}(E|J, M)\mathbf{P}(J|M)\mathbf{P}(M)$

The corresponding BN, which of course is fully connected:



Let us now try to identify the **causal** dependencies among the five variables, that may lead to **conditional independence assumptions** useful to simplify the BN.

To this aim, one can directly build the BN, adding nodes one at a time, with the coresponding edges, without first writing the corresponding expression of the chain rule.

Burglaries and earthquakes can be considered as "root causes". It can also be assumed that there are no causal dependencies among them. Note that this is only a (reasonable) **assumption**, useful to **simplify** the probabilistic model.

Since there is no causal dependency between burglaries and earthquakes, any of the corresponding node can be added first to the BN, e.g., *Burglary*:

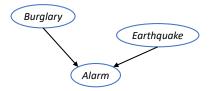


The next node to add is *Earthquake*, with **no** incoming edge from *Burglary*:

Burglary Earthquake

Both burglaries and earthquakes **directly** influence the state of the alarm, but **not** the fact that Mary or John calls: we can assume they will call only if they hear (or believe to hear) the alarm sounding.

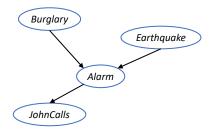
The next node to add is therefore *Alarm*, with incoming edges from both *Burglary* and *Earthquake*:



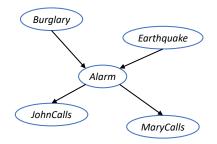
What about JohnCalls and MaryCalls?

- both variables are **directly** influenced by the state of the alarm, not by burglaries and earthquakes
- we can assume John and Mary do not communicate with each other, so neither variable **directly** influences the other

We can therefore add any of these variables, e.g., *JohnCalls*, with an incoming edge only from *Alarm*:

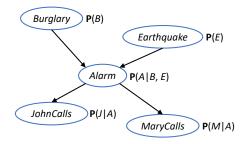


We finally add *MaryCalls*, with an incoming edge only from *Alarm*:



Note that the resulting BN is **not** fully connected as the first one shown above: it contains only 4 edges instead of 10.

It is easy to identify the marginal and conditional distributions associated with each node:



The corresponding expression of the full joint distribution is:

 $\mathbf{P}(B, E, A, J, M) = \mathbf{P}(M|A)\mathbf{P}(J|A)\mathbf{P}(A|E, B)\mathbf{P}(E)\mathbf{P}(B)$

Is the above BN an **exact** and **complete** probabilistic model of the cosidered domain?

It is likely to be only an **approximate** rather than exact model, due to the **assumptions** made. If the assumptions are reasonable, it might be a good approximation.

What about completeness, e.g., other possible causes of the considered events not represented by the BN? For instance

no node for Mary's currently listening to loud music

► no node for the telephone ringing and confusing John Actually all such factors are **implicitly summarised** in the uncertainty associated with the links included in the BN.

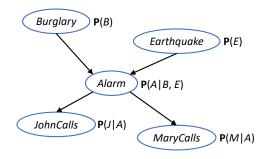
The choice of not representing **all** possible factors **explicitly** is an example of **laziness** and **ignorance**: it would be very difficult to consider them all and evaluate their likelihood.

On the other hand, the distributions associated to a BN **summarise** a **potentially infinite** set of circumstances in which the considered events can happen or not, e.g.:

- ▶ **P**(*A*|*E*, *B*) summarises all the other causes, beside earthquakes and burglaries, that make the alarm sound or fail to sound (passing helicopter, high humidity, power failure, dead battery, cut wires, a dead mouse stuck inside the bell, etc.)
- ▶ **P**(*J*|*A*) and **P**(*M*|*A*) summarises all the other causes, beside the state of the alarm, that make John or Mary call or fail to call (out to lunch, on vacation, listening to loud music, passing helicopter, etc.).

This is the way in which probabilistic models can be kept small enough to be **practically useful**, albeit only **approximate**.

What are the conditional independence assumptions **informally** made when constructing our BN?



To identify conditional independence assumptions, consider

the general expression of the full joint distribution using the chain rule for the chosen order between the variables:

 $\mathbf{P}(B, E, A, J, M) = \mathbf{P}(M|J, A, E, B)\mathbf{P}(J|A, E, B)\mathbf{P}(A|E, B)\mathbf{P}(E|B)\mathbf{P}(B)$

the expression corresponding to the BN:

 $\mathbf{P}(B, E, A, J, M) = \mathbf{P}(M|A)\mathbf{P}(J|A)\mathbf{P}(A|E, B)\mathbf{P}(E)\mathbf{P}(B)$ (cont.)

Comparing the corresponding terms in the two expressions above:

1. Earthquake is (unconditionally) independent on Burglary:

$$\mathbf{P}(E|B) = \mathbf{P}(E)$$

2. JohnCalls is conditionally independent on Burglary and Earthquake given Alarm:

$$\mathbf{P}(J|A, E, B) = \mathbf{P}(J|A)$$

3. *MaryCalls* is conditionally independent on *Burglary*, *Earthquake* and *JohnCalls* given *Alarm*:

$$\mathbf{P}(M|J,A,E,B) = \mathbf{P}(M|A)$$

The ones listed above are the **only** conditional (or absolute) independence assumptions **explicitly** made during the construction of the considered BN.

Nevertheless, it can be shown that a BN also implies other conditional independence assumptions (see later).

Attention should be paid to avoid drawing **wrong** conclusions about independence relations implied by a given BN

 for instance, from the fact that there are no incoming links to Burglary, it would be **not correct** to conclude that Burglary is independent on all the other variables

To define the full joint distribution P(B, E, A, J, M), $2^5 - 1 = 31$ probability values must be specified.

How much do the above conditional independence assumptions simplify its definition?

for a total of 10 probability values, instead of 31.

To construct the above BN the variables have been ordered from "root causes" to "end effects": *Burglary*, *Earthquake*, *Alarm*, *JohnCalls*, *MaryCalls*.

As a consequence, **all** the distributions associated to the nodes of the BN (i.e., to the corresponding expression of the chain rule) are of **causal** form, P(effect|causes), which are usually simpler to estimate, and allow conditional independence relations to be exploited.

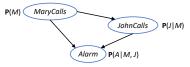
What happens if one choses a different order among the variables to construct a BN?

Consider for instance adding variables in the following order: *MaryCalls*, *JohnCalls*, *Alarm*, *Burglary*, *Earthquake*.

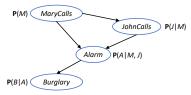
Adding MaryCalls:

Adding JohnCalls: if Mary calls, it is likely the alarm has sounded, which makes it more likely that John calls: JohnCalls needs MaryCalls as a parent

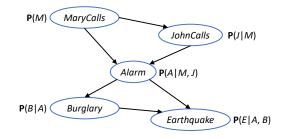
Adding Alarm: if both call, it is more likely that the alarm has sounded than if just one or neither call: both MaryCalls and JohnCalls are needed as parents



Adding Burglary: if we know the alarm is sounding, a call from John or Mary does not provide additional information about burglary: only Alarm is needed as parent



Adding Earthquake: if the alarm is sounding, it is more likely that an earthquake occurred; if we know that also a burglary occurred, this explains the alarm, and the probability of an earthquake would be only slightly above normal; a call from John or Mary does not provide any additional information on earthquakes: only Alarm and Burglary are needed as parents

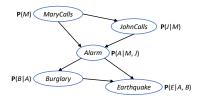


The resulting BN has two more edges than the previous one, and therefore requires **more** probability values to be specified.

Moreover, the required distributions are much difficult to estimate, since they require unnatural probability judgments, and in particular **non-causal** ones, e.g., assessing the probability of an earthquake given that a burglary occurred and the alarm sounded, $\mathbf{P}(E|B, A)$.

Exercise

1. Write the expression of the full joint distribution using the chain rule, corresponding to the BN on the right



- 2. Identify the conditional independence assumptions implicitly made during the construction of the above BN
- 3. Compute the number of probability values to be specified to define the distributions associated to the above BN
- Construct a BN using a different order among the variables: MaryCalls, JohnCalls, Earthquake, Burglary, Alarm, and repeat steps 1–3

Inference in Bayesian networks

Exact inference in Bayesian networks

As previously mentioned, the basic task of probabilistic inference systems is to compute the **posterior** probability distribution of a set of **query** variables \mathbf{Q} given the observed values \mathbf{e} of some distinct set of **evidence** variables \mathbf{E} :

P(Q|E = e)

We have already seen that, in principle, any probabilistic inference can be carried out through the **full** joint distribution of **all** the variables, including the ones that are not part of \mathbf{Q} nor \mathbf{E} , that will be denoted with \mathbf{Y} .

Exact inference in Bayesian networks

For a single query variable Q (all results can be easily extended to more query variables), we known that the posterior distribution $\mathbf{P}(Q|\mathbf{E} = \mathbf{e})$ can be computed through the full joint distribution using the definition of posterior probability and the sum rule:

$$\mathsf{P}(Q|\mathsf{E}=\mathsf{e}) = \frac{\mathsf{P}(Q,\mathsf{E}=\mathsf{e})}{\mathsf{P}(\mathsf{E}=\mathsf{e})} = \frac{\sum_{\mathsf{y}}\mathsf{P}(Q,\mathsf{E}=\mathsf{e},\mathsf{Y}=\mathsf{y})}{\sum_{q,\mathbf{y}}P(Q=q,\mathsf{E}=\mathsf{e},\mathsf{Y}=\mathsf{y})}$$

If the full joint distribution is represented by a BN

- ▶ if E = pa(Q), then P(Q|E = e) = P(Q|pa(Q) = e), which is already known as part of the BN definition
- otherwise, both the numerator and denominator of the above expression can be computed using the expression of the chain rule associated to the BN

Exact inference in Bayesian networks: example

After a call by both Mary and John, one may want to know the posterior probability distribution of a burglary:

$$\mathbf{P}(B|J = true, M = true)$$

Using the definition of conditional probability:

$$\mathbf{P}(B|J = true, M = true) = \frac{\mathbf{P}(B, J = true, M = true)}{P(J = true, M = true)}$$

The numerator and the denominator can be rewritten in terms of the full joint distribution, using the sum rule over all the remaining variables.

(cont.)

Exact inference in Bayesian networks: example

For instance, the numerator can be computed as:

$$\sum_{e,a\in\{true,false\}} \mathbf{P}(B, J = true, M = true, E = e, A = a)$$

In turn, this can be rewritten using the chain rule corresponding to the BN:

$$\sum_{e,a \in \{t,f\}} P(M = t | A = a) P(J = t | A = a) \mathbf{P}(A = a | E = e, B) P(E = e) \mathbf{P}(B)$$

All the conditional and prior distributions in the last expression are known as part of the BN definition.

The denominator, P(J = t, M = t), can be computed similarly.

Computational complexity of exact inference

The **exact** inference procedure shown above has an **exponential** worst-case **time** and **space complexity** in the number of variables.

A practical solution is to trade exactness for a lower complexity.

To this aim, **approximate** inference methods based on **randomised sampling** can be used, known as **Monte Carlo** algorithms, which have become widespread in computer science to estimate quantities difficult to calculate exactly.

In the following, the **direct sampling** family of Monte Carlo algorithms is considered.

More efficient inference methods exist, like the Markov Chain Monte Carlo algorithm.

Direct sampling

Primitive element of sampling algorithms: direct sampling from a known **prior** probability distribution (i.e., with **no** evidence).

Example: the outcome of flipping an unbiased coin can be represented as a random variable $Coin \in \langle heads, tails \rangle$ with prior distribution $P(Coin) = \langle 0.5, 0.5 \rangle$.

Sampling from P(Coin) is equivalent to flipping the coin.

This can be achieved, e.g., by

 generating a random number x with uniform distribution in [0,1] (e.g., using library functions of programming languages, such as Python's random)

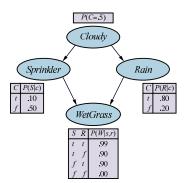
• setting Coin = heads, if x < 0.5, and Coin = tails otherwise

The same method can be used for approximate inference in Bayesian networks.

The sampling process consists of sampling each variable X in topological order, i.e., after its parents pa(X), using the probability distribution conditioned on the values **z** already assigned to its parents, $\mathbf{P}(X|pa(X) = \mathbf{z})$.

Direct sampling: example

A BN that describes a person's daily lawn routine: each morning, she checks the weather; if it is cloudy, she usually does not turn on the sprinkler; if the sprinkler is on, or if it rains during the day, the grass will be wet.



(cont.)

Direct sampling: example

A possible ordering: Cloudy, Sprinkler, Rain, WetGrass

- 1. sample from $P(Cloudy) = \langle 0.5, 0.5 \rangle$ suppose this returns *true*
- 2. sample from P(Sprinkler|Cloudy = t) = (0.1, 0.9)suppose this returns *false*
- 3. sample from $P(Rain|Cloudy = t) = \langle 0.8, 0.2 \rangle$ suppose this returns *true*
- 4. sample from $P(WetGrass|Sprinkler = f, Rain = t) = \langle 0.9, 0.1 \rangle$ suppose this returns *true*

The outcome of the above direct sampling process is therefore:

Cloudy = true, Sprinkler = false, Rain = true, WetGrass = true

Direct sampling: justification

Let $P_{\text{DS}}(X_1 = x_1, \dots, X_n = x_n)$ denote the probability that event (x_1, \dots, x_n) is generated by the direct sampling process from a BN. Looking at the sampling process, it is easy to see that

$$P_{\rm DS}(X_1 = x_1, \dots, X_n = x_n) = \prod_{k=1}^n P(X_k = x_k | pa(X_k))$$

The right-hand side of the above expression equals by definition the full joint distribution of the *n* variables, therefore:

$$P_{\rm DS}(X_1 = x_1, \dots, X_n = x_n) = P(X_1 = x_1, \dots, X_n = x_n)$$
 (cont.)

Direct sampling: justification

If *N* samples are drawn, denoting with $N_{DS}(x_1, \ldots, x_n)$ the number of occurrences of the event (x_1, \ldots, x_n) , we expect its frequency to converge to its probability as *N* increases:

$$\lim_{N \to \infty} \frac{N_{\mathrm{DS}}(x_1, \dots, x_n)}{N} = P_{\mathrm{DS}}(X_1 = x_1, \dots, X_n = x_n)$$
$$= P(X_1 = x_1, \dots, X_n = x_n)$$

For instance, the sampling probability of the event in the previous example is

$$P_{DS}(Cloudy = t, Sprinkler = f, Rain = t, WetGrass = t)$$

$$= P(Cloudy = t) \times P(Sprinkler = f | Cloudy = t)$$

$$\times P(Rain = t | Cloudy = t) \times P(WetGrass = t | Sprinkler = f, Rain = t)$$

$$= 0.5 \times 0.9 \times 0.8 \times 0.9 = 0.324$$

Therefore, for large N we expect 32.4% samples to be of this event.

Rejection sampling in Bayesian networks

Rejection sampling produces samples from a hard-to-sample distribution given an easy-to-sample one.

It can be used to compute a conditional probability $P(X_k = x_k | \mathbf{E} = \mathbf{e})$ from the full joint distribution $\mathbf{P}(X_1, \dots, X_n)$ specified by a BN:

- 1. generate N samples from the prior distribution $\mathbf{P}(X_1, \ldots, X_n)$ using direct sampling
- 2. reject all samples that do not match the evidence
- 3. estimate $P(X_k = x_k | \mathbf{E} = \mathbf{e})$ as the fraction of remaining samples for which $X_k = x_k$

The larger the number of samples, the more accurate the estimate. Its justification is similar to that of direct sampling. Rejection sampling in Bayesian networks: example

Assume one wishes to estimate P(Rain = true | Sprinkler = true)from N = 100 samples.

Among the 100 generated samples, suppose 73 have Sprinkler = true: these samples are rejected.

Among the remaining 27 samples, assume 8 have Rain = true and 19 have Rain = false. Therefore:

$$P(Rain = true | Sprinkler = true) \approx rac{8}{27} \simeq 0.296$$

Likelihood weighting in Bayesian networks

Drawback of rejection sampling: the lower the prior probability of the evidence, $P(\mathbf{E} = \mathbf{e})$, the larger the number of rejected samples before generating enough samples consistent with the evidence to **accurately** estimate $P(X_k = x_k | \mathbf{E} = \mathbf{e})$.

This inefficiency is avoided by **likelihood weighting**, which generates only events that are consistent with the evidence. Informally:

- set the likelihood weight w = 1.0
- ▶ fixing the evidence variables **E** to the observed values **e**
- sampling only the remaining variables X_k and Y
- in the count of the events where X_k = x_k, each one is
 weighted by the likelihood that it accords to the evidence:

$$w \leftarrow w \times P(E_i = e_i | pa(E_i))$$
, for each $E_i \in \mathbf{E}$

Likelihood weighting in Bayesian networks

An example from the previous BN, using again the ordering *Cloudy*, *Sprinkler*, *Rain*, *WetGrass*, to compute

P(*Rain* = *true*|*Sprinkler* = *true*, *Grass* = *true*)

- 1. set the likelihood weight w = 1.0
- 2. sample from $P(Cloudy) = \langle 0.5, 0.5 \rangle$ suppose this returns *true*
- 3. Sprinkler is an evidence variable whose value is true:

$$w \leftarrow w \times P(Sprinkler = true | Cloudy = true) = 0.1$$

- 4. sample from $P(Rain|Cloudy = true) = \langle 0.8, 0.2 \rangle$ suppose this returns *true*
- 5. WetGrass is an evidence variable whose value is true:

 $w \leftarrow w \times P(WetGrass = true | Sprinkler = true, Rain = true) = 0.099$

(cont.)

Likelihood weighting in Bayesian networks

The outcome of the above sampling process is:

Cloudy = true, Sprinkler = true, Rain = true, WetGrass = true,

which is counted among the events for which Rain = true with weight 0.099.

The weight is low since the above event is a cloudy day, which makes the sprinkler unlikely to be on.

Also the justification of the likelihood weighting algorithm is similar to that of direct sampling.