#### **Artificial Intelligence**

Bayesian networks Chapter 14, AIMA

#### Inference

• Inference in the statistical setting means computing probabilities of different outcomes given the observed information

## **P**(*Outcome* | *Information*)

• We need an efficient method for doing this

#### Conditional independence

We say that X and Y are conditionally independent if



What's the relation between independence and conditional independence?

#### Naive Bayes: Combining evidence

Assume full conditional independence and express the full joint probability distribution as:

 $\mathbf{P}(Effect_{1}, Effect_{2}, \dots, Effect_{n}, Cause) = \mathbf{P}(Effect_{1}, Effect_{2}, \dots, Effect_{n} | Cause)\mathbf{P}(Cause) = \mathbf{P}(Effect_{1} | Cause) \cdots \mathbf{P}(Effect_{n} | Cause)\mathbf{P}(cause) = \left[\prod_{i=1}^{n} \mathbf{P}(Effect_{i} | Cause)\right]\mathbf{P}(Cause)$ 

#### Naive Bayes: Dentist example

 $\mathbf{P}(Toothache, Catch, Cavity) =$   $\mathbf{P}(Toothache, Catch | Cavity) \mathbf{P}(Cavity) \approx$   $\mathbf{P}(Toothache | Cavity) \mathbf{P}(Catch | Cavity) \mathbf{P}(Cavity)$   $\Rightarrow \mathbf{P}(toothache, catch, cavity) \approx$   $\frac{(0.108 + 0.012)}{0.2} \times \frac{(0.108 + 0.072)}{0.2} \times 0.2 = 0.108$ 

True value: P(toothache, catch, cavity) = 0.108



	tooth	lache	−toothache		
	catch	$\neg$ catch	catch	$\neg$ catch	
cavity	0.108	0.012	0.072	0.008	
−cavity	0.016	0.064	0.144	0.576	

## Naive Bayes: Dentist example

			P(Toothache Cavity)	toothache	_	toothache		
P(Catch Cavity)	catch	−catch	cavity	0.6		0.4		
cavity	0.9	0.1	−cavity	0.1		0.9		
¬cavity	0.2	0.8	2 independer	nt numbers			P(Cavity)	
2 indeper	ndent num	nbers				cavity	0.2	
						−cavity	0.8	
			_		1	independe	ent numb	er



P(Catch, Toothache,Cavity)	tooth	lache	¬toothache		
	catch	−catch	catch	−catch	
cavity	0.108	0.012	0.072	0.008	
−cavity	0.016	0.064	0.144	0.576	

Full table has  $2^{3}-1=7$  independent numbers  $[O(2^{n})]$ 

#### Naive Bayes application: Learning to classify text

Use a dictionary with words (not too frequent and not too infrequent), e.g.
 w<sub>1</sub> = airplane, w<sub>2</sub> = algorithm, ...



- Estimate conditional probabilities
   P(w<sub>i</sub> | interesting) and P(w<sub>i</sub> | uninteresting)
- Compute P(text | interesting) and P(text | uninteresting) using Naive Bayes
   (and assuming that word position in text is unimportant)

$$P(\text{text} | \text{interesting}) \approx \prod_{i} P(w_i | \text{interesting})$$

Where  $w_i$  are the words occuring in this particular text.

#### Naive Bayes application: Learning to classify text

• Then compute the probability that the text is interesting (or uninteresting) using Bayes' theorem

 $P(\text{interesting} | \text{text}) = \frac{P(\text{interesting})P(\text{text} | \text{interesting})}{P(\text{text})}$ 

P(text) is just a normalization factor; it is not necessary to compute it since we are only interested in knowing whether

P(interesting | text) > P(uninteresting | text)



#### Inference

• Inference in the statistical setting means computing probabilities of different outcomes given the observed information

## **P**(*Outcome* | *Information*)

• We need an efficient method for doing this which is more widely applicable than the naïve Bayes model

#### Bayesian networks

- A **Bayesian network** is a directed graph in which each node is annotated with quantitative probability information:
- 1. The set of nodes of the network corresponds to a set of random variables  $\{X_1, X_2, X_3, ...\}$
- 2. pairs of nodes can be connected by directed links defining a parent  $\rightarrow$  child relation
- 3. Each node  $X_i$  contains a conditional probability distribution  $P(X_i | Parents(X_i))$
- 4. The graph is a directed acyclic graph (DAG)



	tooth	lache	¬toothache		
	catch	¬catch	catch	¬catch	
cavity	0.108	0.012	0.072	0.008	
¬cavity	0.016	0.064	0.144	0.576	

#### The alarm network



Burglar alarm responds to both earthquakes and burglars.

Two neighbors: John and Mary, who have promised to call you when the alarm goes off.

John always calls when there's an alarm, and sometimes when there's not an alarm.

Mary sometimes misses the alarms (she likes loud music).

#### The cancer network





### Meaning of Bayesian network

The general chain rule (always true):

$$P(x_1, x_2, \dots, x_n) = P(x_1 | x_2, x_3, \dots, x_n) P(x_2, x_3, \dots, x_n) = P(x_1 | x_2, x_3, \dots, x_n) P(x_2 | x_3, x_4, \dots, x_n) P(x_3, x_4, \dots, x_n) = \cdots$$
$$= \prod_{i=1}^n P(x_i | x_{i+1}, \dots, x_n)$$

The Bayesian network chain rule:

$$P(x_1, x_2, \dots, x_n) = \prod_{i=1}^n P(x_i \mid parents(X_i))$$

The BN is a correct representation of the domain <u>iff</u> each node is conditionally independent of its predecessors, given its parents.

#### Bayes network node is a function



#### Bayes network node is a function



Can be any type of function from values to distributions.



*Note: Each number in the tables represents a boolean distribution.* 

*Hence, for every input, there is a distribution as an output.* 



#### The alarm network



*The fully correct alarm network might look something like the figure.* 

The Bayesian network assumes that some of the variables are independent • or that the dependencies can be neglected since they are very weak

The correctness of the Bayesian network, of course, depends on the validity of these assumptions!

*It is this sparse connection structure that makes the BN approach feasible: ~linear growth in complexity rather than exponential, in practice* 

#### How to construct a BN?

• Add nodes in causal order

"causal" determined from expertise

- Determine conditional independence using either (or all) of the following semantics:
  - Blocking/d-separation rule
  - Non-descendant rule
  - Markov blanket rule
  - Experience/your beliefs

#### Path blocking & d-separation



Intuitively, knowledge about Serum Calcium influences our belief about Cancer – if we don't know the value of Cancer – which, in turn, influences our belief about Lung Tumour, etc.

However, if we are given the value of Cancer (i.e. C= true or false), then knowledge of Serum Calcium will not tell us anything about Lung Tumour that we don't already know

- this is conditional independence at work!

We say that Cancer **d-separates** (direction-dependent separates) Serum Calcium and Lung Tumour.

## Path blocking & d-separation

- $X_i$  and  $X_j$  are d-separated if all paths betweeen them are blocked Two nodes  $X_i$  and  $X_j$  are conditionally independent given a set  $\Omega = \{X_1, X_2, X_3, ...\}$  of nodes if for every undirected path in the BN between  $X_i$  and  $X_j$  there is some node  $X_k$  on the path having one of the following three properties:
- $\begin{array}{ll} 1. & X_k \in \varOmega \text{, and both arcs on the path} \\ \text{ lead out of } X_k \text{.} \end{array} \end{array}$
- 2.  $X_k \in \Omega$ , and one arc on the path leads into  $X_k$  and one arc leads out.
- 3. Neither  $X_k$  nor any descendant of  $X_k$  is in  $\Omega$ , and both arcs on the path lead into  $X_k$ .
- $X_k$  **blocks** the path between  $X_i$  and  $X_j$



 $P(X_i, X_j \mid \Omega) = P(X_i \mid \Omega) P(X_j \mid \Omega)$ 

#### Some definitions of BN (from Wikipedia)

**1.** X is a Bayesian network with respect to G if its joint probability density function can be written as a product of the individual density functions, conditional on their parent variables:

$$P(x_1, x_2, \dots, x_n) = \prod_{i=1}^n P(x_i \mid parents(X_i))$$

 $\mathbf{X} = \{X_1, X_2, ..., X_N\}$  is a set of random variables  $\mathbf{G} = (V, E)$  is a directed acyclic graph (DAG) of vertices (V) and edges (E)

#### Some definitions of BN (from Wikipedia)

**1.** X is a Bayesian network with respect to G if it satisfies the local Markov property: each variable is conditionally independent of its non-descendants given its parent variables:

$$P(x_1 \mid non - descentants(X_i)) = P(x_i \mid parents(X_i))$$

Note:  $parents(X_i) \subseteq non - descentants(X_i)$ 

 $X = \{X_1, X_2, ..., X_N\}$  is a set of random variables G = (V, E) is a directed acyclic graph (DAG) of vertices (V) and edges (E)

#### Non-descendants

Um

Yn

Z<sub>ni</sub>

Х

Z<sub>1j</sub>

 $Y_1$ 



$$P(X, Z_{1j}, ..., Z_{nj} | U_1, ..., U_m) =$$

$$P(X | U_1, ..., U_m) P(Z_{1j}, ..., Z_{nj} | U_1, ..., U_m)$$

$$P(X | Z_{1j}, ..., Z_{nj}, U_1, ..., U_m) =$$

$$P(X | U_1, ..., U_m)$$

#### Some definitions of BN (from Wikipedia)

 X is a Bayesian network with respect to G if every node is conditionally independent of all other nodes in the network, given its Markov blanket.

The Markov blanket of a node is its parents, children and children's parents.

 $P(x_i | \text{all nodes}) = P(x_i | Markov blanket(X_i))$ 

 $X = \{X_1, X_2, ..., X_N\}$  is a set of random variables G = (V, E) is a directed acyclic graph (DAG) of vertices (V) and edges (E)

## Markov blanket

 $X_3$ 

 $X_{4}$ 

 $X_2$ 

 $X_1$ A node is conditionally independent of all U₁  $U_{m}$ other nodes in the network, given its parents, children, and Х  $Z_{1j}$ Z<sub>nj</sub> children's parents These constitute the node's Markov Y<sub>1</sub> Yn blanket, X<sub>5</sub>  $P(X | X_1, ..., X_k, U_1, ..., U_m, Z_{1i}, ..., Z_{ni}, Y_1, ..., Y_n) =$  $X_6$  $P(X | U_1, ..., U_m, Z_{1i}, ..., Z_{ni}, Y_1, ..., Y_n)$  $X_k$  $P(X, X_1, ..., X_k | U_1, ..., U_m, Z_{1i}, ..., Z_{ni}, Y_1, ..., Y_n) =$  $P(X | U_1, ..., U_m, Z_{1i}, ..., Z_{ni}, Y_1, ..., Y_n) P(X_1, ..., X_k | U_1, ..., U_m, Z_{1i}, ..., Z_{ni}, Y_1, ..., Y_n)$ 

#### Some definitions of BN (from Wikipedia)

**1.** X is a Bayesian network with respect toG if, for any two nodes i, j:

$$P(x_i, x_j | X_1, X_2, ..., X_N) =$$

 $P(x_i | d - separating set(i, j))P(x_j | d - separating set(i, j))$ 

The *d*-separating set(*i*,*j*) is the set of nodes that *d*-separate node *i* and *j*.

The Markov blanket of node *i* is the minimal set of nodes that d-separates node *i* from all other nodes.

 $\mathbf{X} = \{X_1, X_2, ..., X_N\}$  is a set of random variables  $\mathbf{G} = (V, E)$  is a directed acyclic graph (DAG) of vertices (V) and edges (E)

#### Causal networks

- Bayesian networks are usually used to represent causal relationships. This is, however, not strictly <u>necessary</u>: a directed edge from node i to node j does not require that X<sub>i</sub> is causally dependent on X<sub>i</sub>.
  - This is demonstrated by the fact that Bayesian networks on the two graphs:



*are equivalent. They impose the same conditional independence requirements.* 

A <u>causal network</u> is a Bayesian network with an explicit requirement that the relationships be causal.

#### Causal networks

$$P(A, B, C) = P(A | B)P(B | C)P(C)$$
  
=  $P(A | B) * \dots * \dots$ 



The equivalence is proved with Bayes theorem...

- Two astronomers in different parts of the world make measurements M<sub>1</sub> and M<sub>2</sub> of the number of stars N in some small region of the sky, using their telescopes. Normally there is a small possibility e of error up to one star in each direction. Each telescope can also (with a much smaller probability f) be badly out of focus (events F<sub>1</sub> and F<sub>2</sub>) in which case the scientist will undercount by three or more stars (or, if N is less than 3, fail to detect any stars at all). Consider the three networks in Figure 14.22<sup>\*</sup>.
  - (a) Which of these Bayesian networks are correct (but not necessarily efficient) representations of the preceeding information?

\*In the 2:nd edition is this exercise 14.3 and the figure is 14.19.

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  - (a) Which of these Bayesian networks are correct (but not necessarily efficient) representations of the preceeding information?

 $F_1$ 



 $M_2$ 



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  - (a) Which of these Bayesian networks are correct (but not necessarily efficient) representations of the preceeding information?



 (i) must be incorrect – N is d-separated from F<sub>1</sub> (or F<sub>2</sub>), relative to {M<sub>1</sub>} (or {M<sub>2</sub>}) i.e. knowing the focus states F would not affect N if we know M:

This cannot be correct!



- Two astronomers in different parts of the world make measurements M<sub>1</sub> and M<sub>2</sub> of the number of stars N in some small region of the sky, using their telescopes. Normally there is a small possibility e of error up to one star in each direction. Each telescope can also (with a much smaller probability f) be badly out of focus (events F<sub>1</sub> and F<sub>2</sub>) in which case the scientist will undercount by three or more stars (or, if N is less than 3, fail to detect any stars at all). Consider the three networks in Figure 14.22.
  - (a) Which of these Bayesian networks are correct (but not necessarily efficient) representations of the preceeding information?



• (*ii*) is correct – *it describes the causal relationships. It is a causal network.* 



- Two astronomers in different parts of the world make measurements M<sub>1</sub> and M<sub>2</sub> of the number of stars N in some small region of the sky, using their telescopes. Normally there is a small possibility e of error up to one star in each direction. Each telescope can also (with a much smaller probability f) be badly out of focus (events F<sub>1</sub> and F<sub>2</sub>) in which case the scientist will undercount by three or more stars (or, if N is less than 3, fail to detect any stars at all). Consider the three networks in Figure 14.22.
  - (a) Which of these Bayesian networks are correct (but not necessarily efficient) representations of the preceeding information?



 (iii) is also ok – a fully connected graph would be correct (but not efficient). (iii) has all connections except M<sub>i</sub>-F<sub>j</sub> and F<sub>i</sub>-F<sub>j</sub>. (iii) is not causal and not efficient.



(*ii*) says: P(F1, F2, N, M1, M2) = P(F1)P(F2)P(N)P(M1 | F1, N)P(M2 | F2, N)

(iii) says:

P(F1, F2, N, M1, M2) = P(M1)P(F1 | N, M1)P(M2 | M1)P(N | M1, M2)P(F2 | N, M2)



(ii) says: P(F1, F2, N, M1, M2) = P(F1)P(F2)P(N)P(M1 | F1, N)P(M2 | F2, N)

 $\begin{array}{l} \text{The full correct expression (one version) is:} \\ P(F1,F2,N,M1,M2) = P(M1 \mid F1,F2,N,M2)P(F1,F2,N,M2) = \\ P(M1 \mid F1,F2,N,M2)P(M2 \mid F1,F2,N)P(F1,F2,N) = \\ P(M1 \mid F1,F2,N,M2)P(M2 \mid F1,F2,N)P(F1 \mid F2,N)P(F2,N) = \\ P(M1 \mid F1,F2,N,M2)P(M2 \mid F1,F2,N)P(F1 \mid F2,N)P(F2 \mid N)P(N) \\ \approx P(M1 \mid F1,N) \\ \approx P(M2 \mid F2,N) \\ \approx P(F1) \\ \approx P(F2) \\ \begin{array}{l} \text{ok} \end{array}$ 

*This is not as efficient as (ii). This requires more conditional probabilities.* 

(iii) says:

P(F1, F2, N, M1, M2) = P(M1)P(F1 | N, M1)P(M2 | M1)P(N | M1, M2)P(F2 | N, M2)

The full correct expression (another version) is: P(F1, F2, N, M1, M2) = P(F1 | F2, N, M1, M2)P(F2, N, M1, M2) = P(F1 | F2, N, M1, M2)P(F2 | N, M1, M2)P(N, M1, M2) = P(F1 | F2, N, M1, M2)P(F2 | N, M1, M2)P(N | M1, M2)P(M1, M2) = P(F1 | F2, N, M1, M2)P(F2 | N, M1, M2)P(N | M1, M2)P(M2 | M1)P(M1) $\approx P(F1 | N, M1) \approx P(F2 | N, M2)$ ok

(i) says:

P(F1, F2, N, M1, M2) = P(F1)P(F2)P(M1 | F1)P(M2 | F2)P(N | M1, M2)



(i) says:

P(F1, F2, N, M1, M2) = P(F1)P(F2)P(M1 | F1)P(M2 | F2)P(N | M1, M2)

 $\begin{array}{l} \text{The full correct expression (a third version) is:} \\ P(F1,F2,N,M1,M2) = P(N \mid M1,M2,F1,F2)P(M1,M2,F1,F2) = \\ P(N \mid M1,M2,F1,F2)P(M1 \mid M2,F1,F2)P(M2,F1,F2) = \\ P(N \mid M1,M2,F1,F2)P(M1 \mid M2,F1,F2)P(M2 \mid F1,F2)P(F1,F2) = \\ P(N \mid M1,M2,F1,F2)P(M1 \mid M2,F1,F2)P(M2 \mid F1,F2)P(F1 \mid F2)P(F2) \\ \approx P(N \mid M1,M2) \\ \approx P(M1 \mid F1) \\ \approx P(M2 \mid F2) \\ \approx P(F1) \end{array}$ 

(i) says: P(F1, F2, N, M1, M2) = P(F1)P(F2)P(M1 | F1)P(M2 | F2)P(N | M1, M2)

This is an unreasonable approximation. The rest is ok.

The full correct expression (a third version) is:

$$\begin{split} P(F1, F2, N, M1, M2) &= P(N \mid M1, M2, F1, F2) P(M1, M2, F1, F2) = \\ P(N \mid M1, M2, F1, F2) P(M1 \mid M2, F1, F2) P(M2, F1, F2) = \\ P(N \mid M1, M2, F1, F2) P(M1 \mid M2, F1, F2) P(M2 \mid F1, F2) P(F1, F2) = \\ P(N \mid M1, M2, F1, F2) P(M1 \mid M2, F1, F2) P(M2 \mid F1, F2) P(F1 \mid F2) P(F2) \\ &\approx P(N \mid M1, M2) \qquad \approx P(M1 \mid F1) \qquad \approx P(M2 \mid F2) \qquad \approx P(F1) \end{split}$$

#### Efficient representation of PDs



- Continuous  $\rightarrow$  Discrete
- Continuous → Continuous

#### Noisy-OR example Boolean → Boolean

$P(E C_1, C_2, C_3)$								
$C_1$	0	1	0	0	1	1	0	1
<i>C</i> <sub>2</sub>	0	0	1	0	1	0	1	1
<i>C</i> <sub>3</sub>	0	0	0	1	0	1	1	1
P(E=0)	1	0.1	0.1	0.1	0.01	0.01	0.01	0.001
P(E=1)	0	0.9	0.9	0.9	0.99	0.99	0.99	0.999

The effect (E) is off (false) when none of the causes are true. The probability for the effect increases with the number of true causes.

$$P(E=0) = 10^{-(\#True)}$$
 (for this example)

Example from L.E. Sucar

#### Noisy-OR general case Boolean → Boolean

$$P(E=0 \mid C_1, C_2, \dots, C_n) = \prod_{i=1}^n q_i^{C_i}$$

$$C_i = \begin{cases} 1 & \text{if true} \\ 0 & \text{if false} \end{cases}$$

Example on previous slide used  $q_i = 0.1$  for all i.

Needs only n parameters, not 2<sup>n</sup> parameters.



Image adapted from Laskey & Mahoney 1999

- Fever *is True if and only if* Cold, Flu *or* Malaria *is True.*
- each cause has an independent chance of causing the effect.
  - all possible causes are listed
  - inhibitors are independent



- $P(\text{Fever} \mid \text{Cold}) = 0.4 \Rightarrow q_1 = 0.6$
- $P(\text{Fever} | \text{Flu}) = 0.8 \Rightarrow q_2 = 0.2$
- $P(\text{Fever} \mid \text{Malaria}) = 0.9 \Rightarrow q_3 = 0.1$



- $P(\text{Fever} \mid \text{Cold}) = 0.4 \Rightarrow q_1 = 0.6$
- $P(\text{Fever} | \text{Flu}) = 0.8 \Rightarrow q_2 = 0.2$
- $P(\text{Fever} \mid \text{Malaria}) = 0.9 \Rightarrow q_3 = 0.1$

 $P(\neg Fever \mid \neg Cold, \neg Flu, \neg Malaria) = 0.6^{\circ} \times 0.2^{\circ} \times 0.1^{\circ} = 1$  $P(Fever \mid \neg Cold, \neg Flu, \neg Malaria) = 1 - 1 = 0$ 



- $P(\text{Fever} \mid \text{Cold}) = 0.4 \Rightarrow q_1 = 0.6$
- $P(\text{Fever} | \text{Flu}) = 0.8 \Rightarrow q_2 = 0.2$
- $P(\text{Fever} \mid \text{Malaria}) = 0.9 \Rightarrow q_3 = 0.1$

 $P(\neg Fever | \neg Cold, \neg Flu, Malaria) = 0.6^{\circ} \times 0.2^{\circ} \times 0.1^{\circ} = 0.1$  $P(Fever | \neg Cold, \neg Flu, Malaria) = 1 - 0.1 = 0.9$ 



- $P(\text{Fever} \mid \text{Cold}) = 0.4 \Rightarrow q_1 = 0.6$
- $P(\text{Fever} | \text{Flu}) = 0.8 \Rightarrow q_2 = 0.2$
- $P(\text{Fever} \mid \text{Malaria}) = 0.9 \Rightarrow q_3 = 0.1$

 $P(\neg Fever | Cold, Flu, \neg Malaria) = 0.6^{1} \times 0.2^{1} \times 0.1^{0} = 0.12$  $P(Fever | Cold, Flu, \neg Malaria) = 1 - 0.12 = 0.88$ 



#### Parametric probability densities Boolean/Discr./Continuous → Continuous

Use parametric probability densities, e.g., the normal distribution

$$P(X) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left(\frac{-(x-\mu)^2}{2\sigma^2}\right) = N(\mu,\sigma)$$

Gaussian networks (a = input to the node)

$$P(X) = \frac{1}{\sigma\sqrt{2\pi}} \exp\left[\frac{-(x-\alpha-\beta a)^2}{2\sigma^2}\right]$$

#### Normal Distribution



Probit & Logit Discrete → Boolean

# *If the input is continuous but output is boolean, use probit or logit*

Logit: 
$$P(A = a | x) = \frac{1}{1 + \exp[-2(\mu - x)/\sigma]}$$
  
Probit:  $P(A = a | x) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{x} \exp(-(x - \mu)^2 / \sigma^2) dx$ 



#### The cancer network



#### Inference in BN

Inference means computing P(X|e), where X is a query (variable) and e is a set of evidence variables (for which we know the values).

Examples:

P(Burglary | john\_calls, mary\_calls)
P(Cancer | age, gender, smoking, serum\_calcium)
P(Cavity | toothache, catch)

#### Exact inference in BN

$$\mathbf{P}(X \mid \mathbf{e}) = \frac{\mathbf{P}(X, \mathbf{e})}{\mathbf{P}(\mathbf{e})} = \alpha \mathbf{P}(X, \mathbf{e}) = \alpha \sum_{\mathbf{y}} \mathbf{P}(X, \mathbf{e}, \mathbf{y})$$

"Doable" for boolean variables: Look up entries in conditional probability tables (CPTs).













A lot of unneccesary repeated computation...

## Complexity of exact inference

- By eliminating repeated calculation & uninteresting paths we can speed up the inference a lot.
- Linear time complexity for singly connected networks (polytrees).
- Exponential for multiply connected networks.
  - Clustering can improve this

### Approximate inference in BN

- Exact inference is intractable in large multiply connected BNs ⇒ use approximate inference: Monte Carlo methods (random sampling).
  - Direct sampling
  - Rejection sampling
  - Likelihood weighting
  - Markov chain Monte Carlo

## Markov chain Monte Carlo

- 1. Fix the evidence variables  $(E_1, E_2, ...)$  at their given values.
- 2. Initialize the network with values for all other variables, including the query variable.
- 3. Repeat the following many, many, many times:
  - a. Pick a non-evidence variable at random (query  $X_i$  or hidden  $Y_j$ )
  - b. Select a new value for this variable, conditioned on the current values in the variable's Markov blanket.

Monitor the values of the query variables.