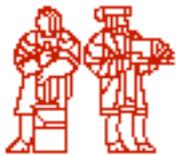


Bayes Networks

6.872/HST.950



What Probabilistic Models Should We Use?

- Full joint distribution
 - Completely expressive
 - Hugely data-hungry
 - Exponential computational complexity
- Naive Bayes (full conditional independence)
 - Relatively concise
 - Need data $\sim (\#\text{hypotheses}) \times (\#\text{features}) \times (\#\text{feature-vals})$
 - Fast $\sim (\#\text{features})$
 - Cannot express dependencies among features or among hypotheses
 - Cannot consider possibility of multiple hypotheses co-occurring

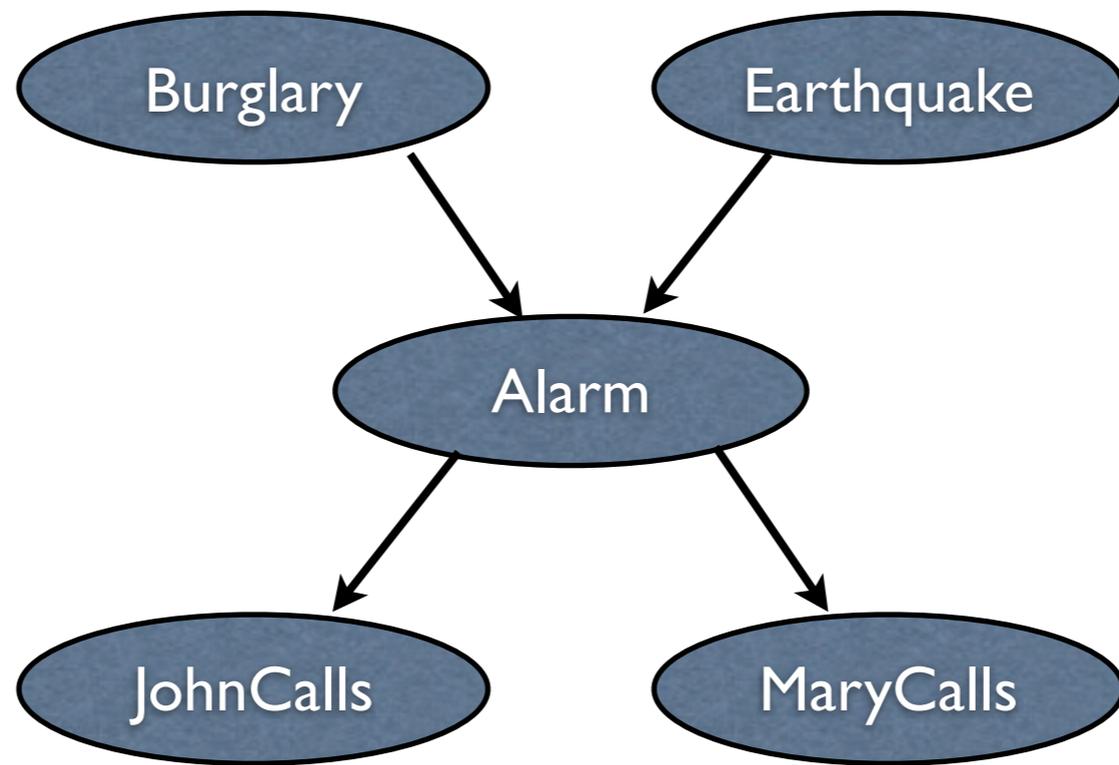
Bayesian Networks

(aka Belief Networks)

- Graphical representation of dependencies among a set of random variables
 - Nodes: variables
 - Directed links to a node from its *parents*: direct probabilistic dependencies
 - Each X_i has a conditional probability distribution, $P(X_i|\text{Parents}(X_i))$, showing the effects of the parents on the node.
 - The graph is directed (DAG); hence, no cycles.
- This is a language that can express dependencies between Naive Bayes and the full joint distribution, more concisely
 - Given some new evidence, how does this affect the probability of some other node(s)? $P(X|E)$ —*belief propagation/updating*
 - Given some evidence, what are the most likely values of other variables? $\text{argmax}_{\mathbf{X}} P(\mathbf{X}|E)$ —*MAP explanation*

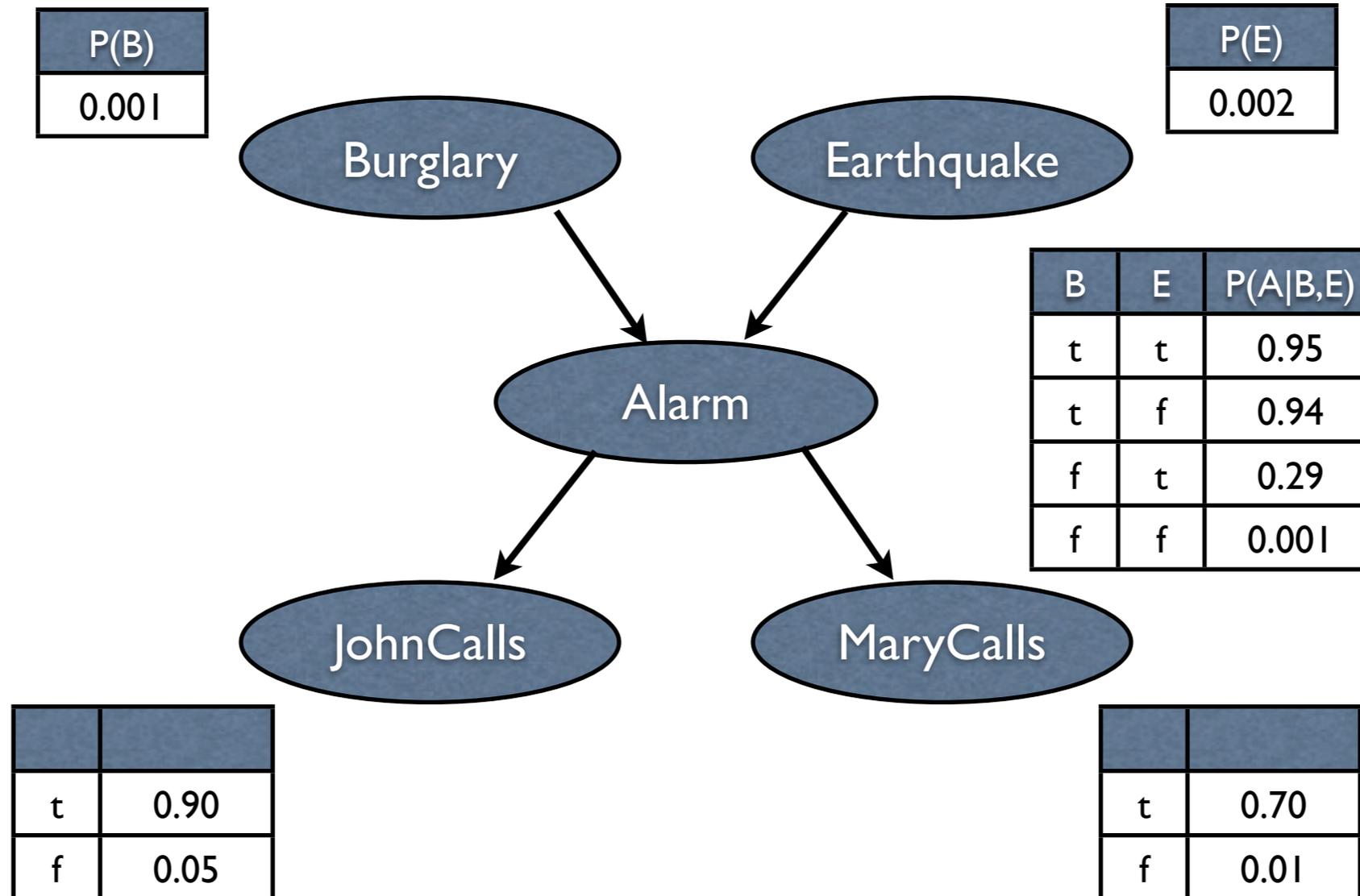
Burglary Network

(due to J. Pearl)

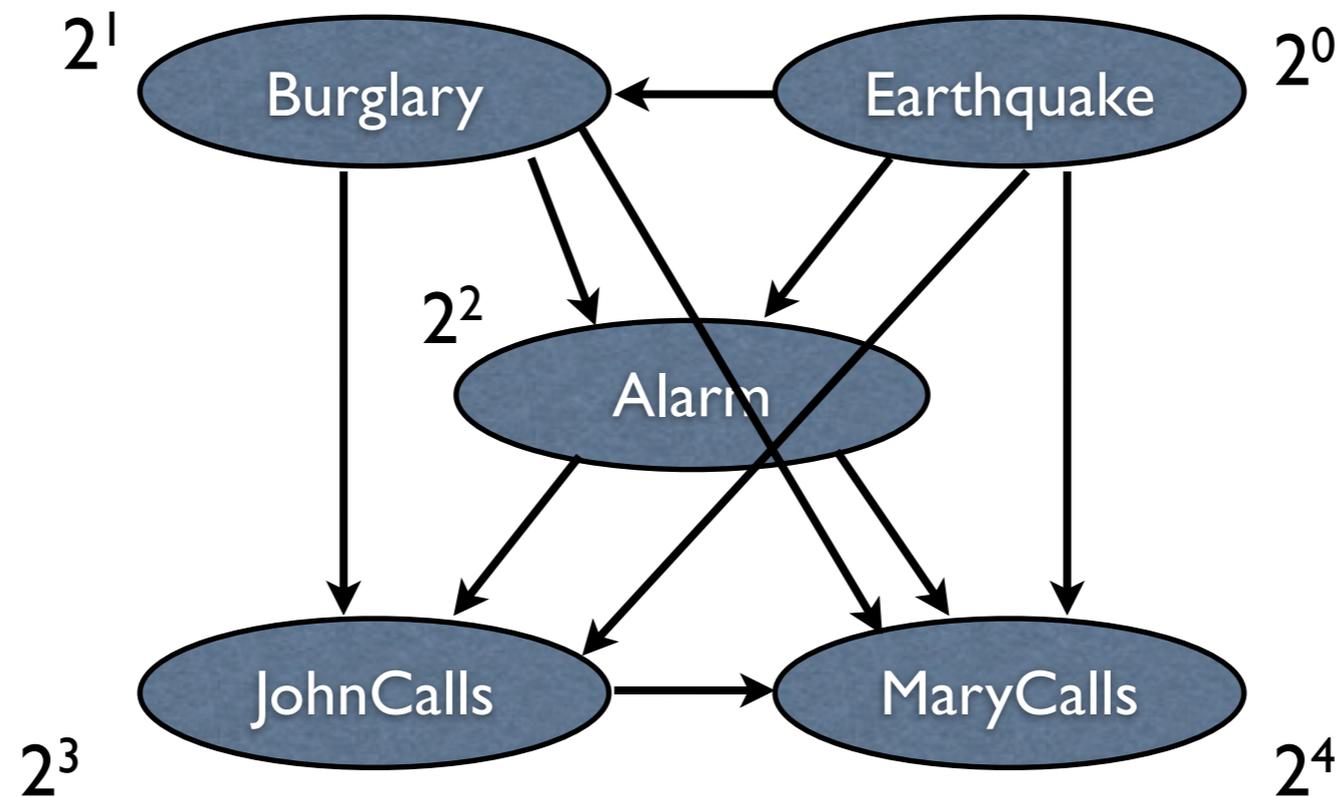


Burglary Network

(due to J. Pearl)



If everything depends on everything



- This model requires just as many parameters as the full joint distribution!

Computing the Joint Distribution from a Bayes Network

- As usual, we abuse notation:

$P(X_1 = x_1 \wedge \dots \wedge X_n = x_n)$ is written as $P(x_1, \dots, x_n)$

- $$P(x_1, \dots, x_n) = \prod_{i=1}^n P(x_i | \text{Par}(X_i))$$

- E.g., what's the probability that an alarm has sounded, there was neither an earthquake nor a burglary, and both John and Mary called?

$$\begin{aligned} & P(j \wedge m \wedge a \wedge \neg b \wedge \neg e) \\ &= P(J|a)P(m|a)P(a|\neg b \wedge \neg e)P(\neg e)P(\neg b) \\ &= 0.90 \times 0.70 \times 0.001 \times 0.999 \times 0.998 = 0.00062 \end{aligned}$$

Requirements for Constructing a BN

- Recall that the definition of the conditional probability was

$$P(x|y) = P(x \wedge y)/P(y)$$

- and thus we get the *chain rule*,

$$P(x \wedge y) = P(x|y)P(y)$$

- Generalizing to n variables,

$$P(x_1, \dots, x_n) = P(x_n|x_{n-1}, \dots, x_1)P(x_{n-1}, \dots, x_1)$$

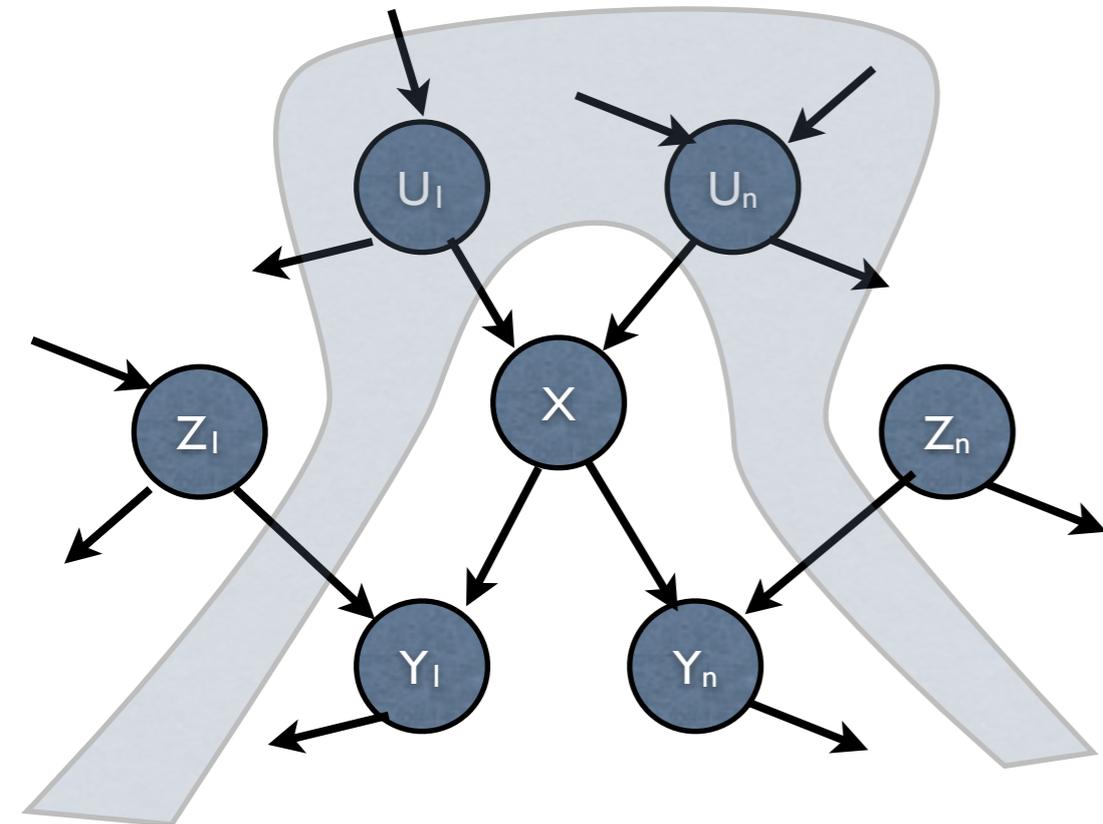
- and repeatedly applying this idea,

$$\begin{aligned} P(x_1, \dots, x_n) &= P(x_n|x_{n-1}, \dots, x_1)P(x_{n-1}|x_{n-2}, \dots, x_1) \cdots P(x_2|x_1)P(x_1) \\ &= \prod_{i=1}^n P(x_i|x_{i-1}, \dots, x_1) \\ &= \prod_{i=1}^n P(x_i|\text{Par}(x_i)) \end{aligned}$$

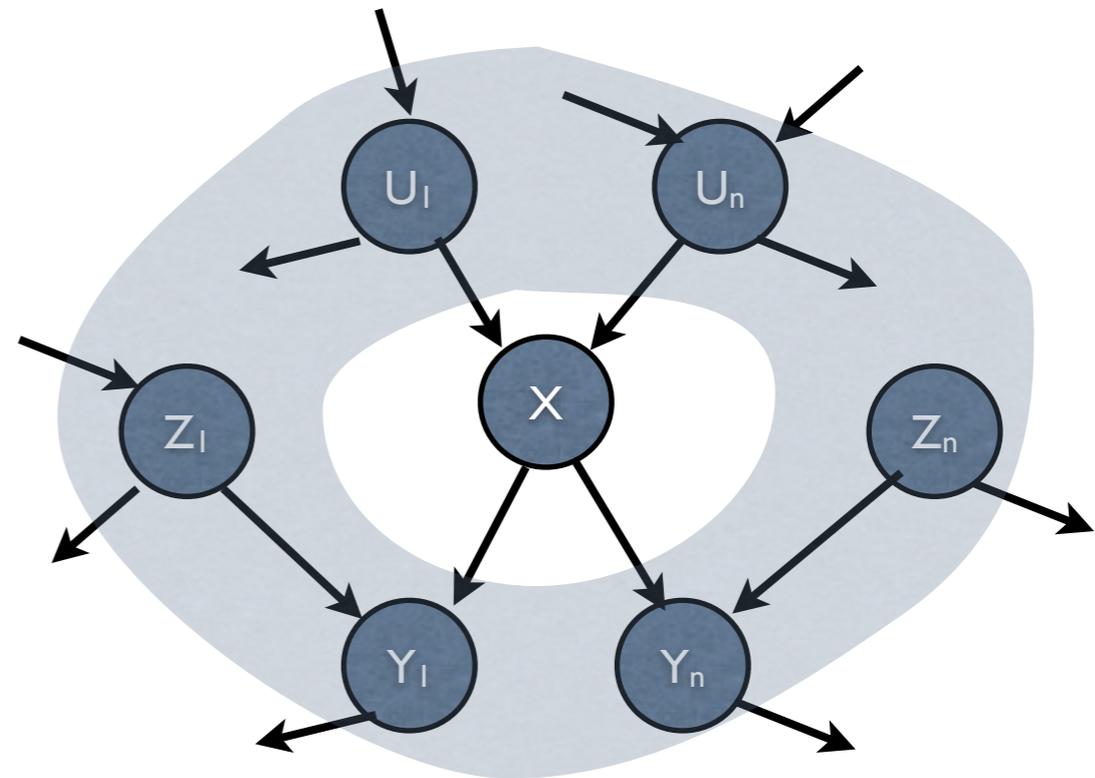
- This “works” just in case we can define a partial order so that

$$\text{Par}(X_i) \subseteq \{X_{i-1}, \dots, X_1\}$$

Topological Interpretations



A node, X , is *conditionally independent* of its non-descendants, Z_i , given its parents, U_i .

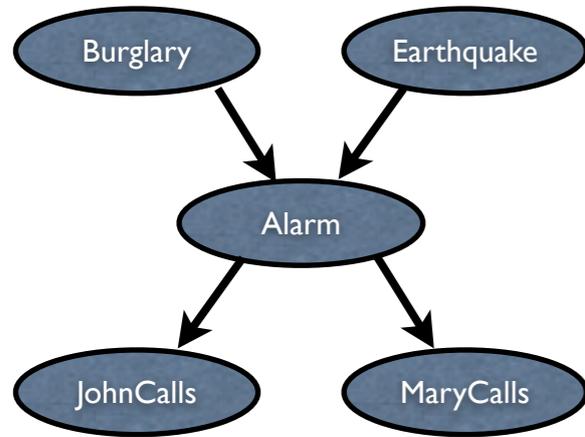


A node, X , is *conditionally independent* of all other nodes in the network given its *Markov blanket*: its parents, U_i , children, Y_i , and children's parents, Z_i .

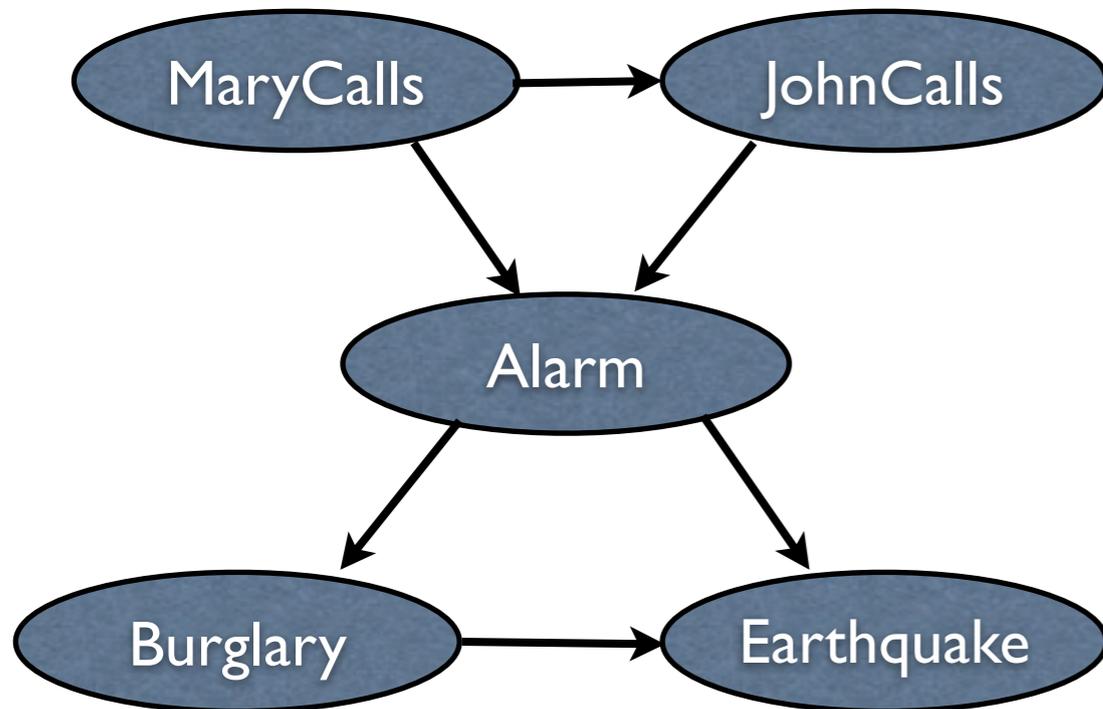
BN's can be Compact

- For a network of 40 binary variables, the full joint distribution has 2^{40} entries ($> 1,000,000,000,000$)
- If $|\text{Par}(x_i)| \leq 5$, however, then the 40 (conditional) probability tables each have ≤ 32 entries, so the total number of parameters $\leq 1,280$
- Largest medical BN I know (Pathfinder) had 109 variables! $2^{109} \approx 10^{36}$

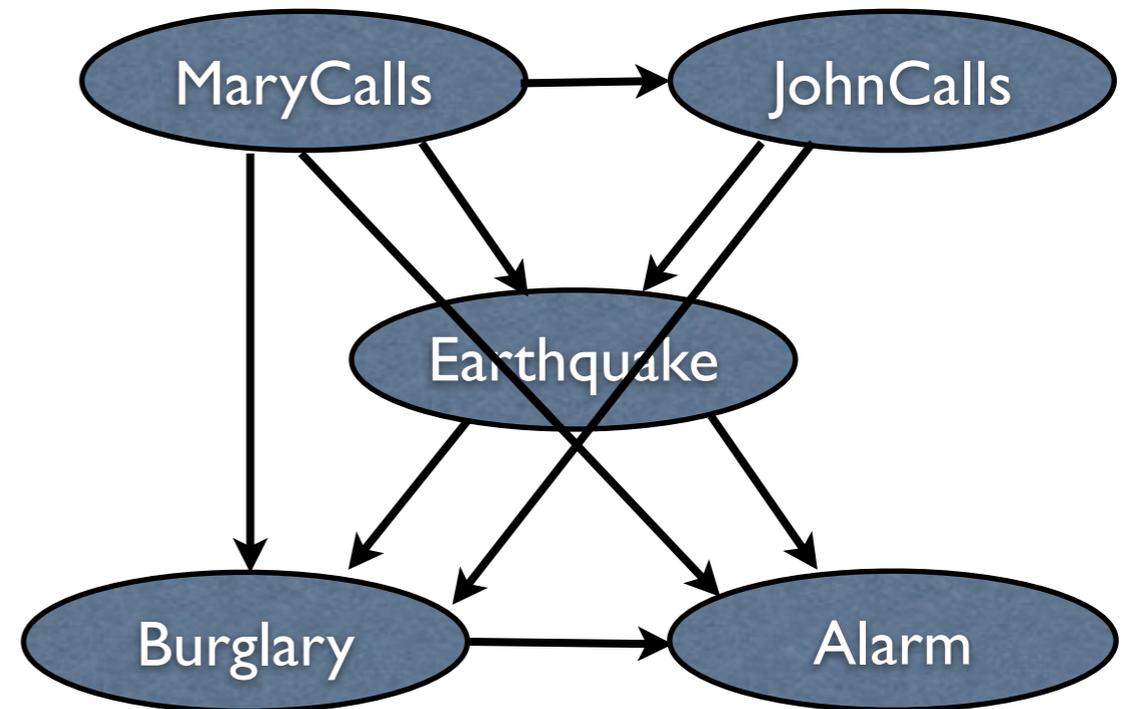
How *Not* to Build BN's



- With the wrong ordering of nodes, the network becomes more complicated, and requires more (and more difficult) conditional probability assessments



Order: M, J, A, B, E



Order: M, J, E, B, A

Simplifying Conditional Probability Tables

- Do we know any structure in the way that $\text{Par}(x)$ “cause” x ?
- If each destroyer can sink the ship with probability $P(s|d_i)$, what is the probability that the ship will sink if it’s attacked by both?
 $(1 - P(s|d_1, d_2)) = (1 - P(s|d_1))(1 - P(s|d_2))$ $(1 - l)$
- For $|\text{Par}(x)| = n$, this requires $O(n)$ parameters, not $O(k^n)$



Image by MIT OpenCourseWare.

d_1



Photo by Konabish on Flickr.

s

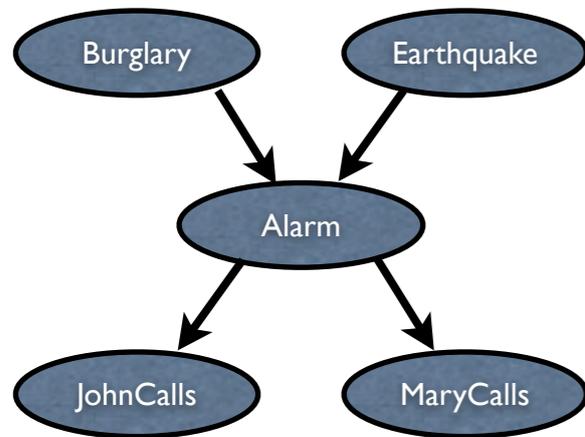


Image by MIT OpenCourseWare.

d_2

Inference

- Recall the two basic inference problems: Belief propagation & MAP explanation
 - Trivially, we can enumerate all “matching” rows of the joint probability distribution
 - For *poly-trees* (not even undirected loops—i.e., only one connection between any pair of nodes; like our Burglary example), there are efficient linear algorithms, similar to constraint propagation
 - For arbitrary BN’s, all inference is NP-hard!
 - Exact solutions
 - Approximation



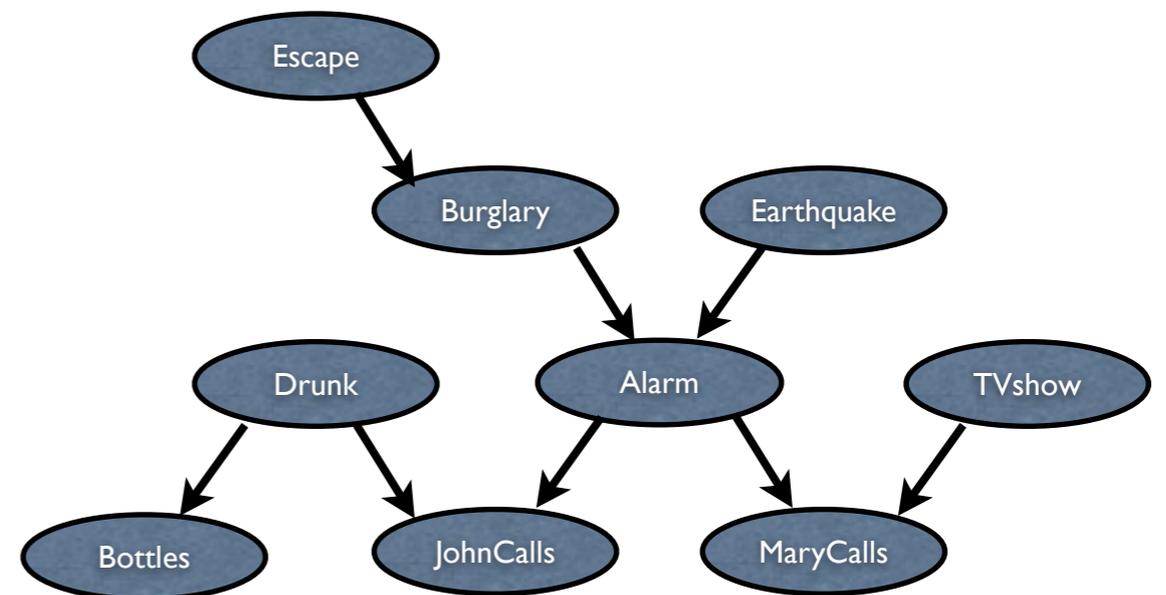
Exact Solution of BN's (Burglary example)

$$\begin{aligned}
 P(b|j, m) &= \alpha \sum_e \sum_a P(b)P(e)P(a|b, e)P(j|a)P(m|a) \\
 &= \alpha P(b) \sum_e P(e) \sum_a P(a|b, e)P(j|a)P(m|a) \\
 P(\mathbf{B}|j, m) &= \alpha\{0.00059224, 0.0014919\} \approx \{0.28, 0.72\}
 \end{aligned}$$

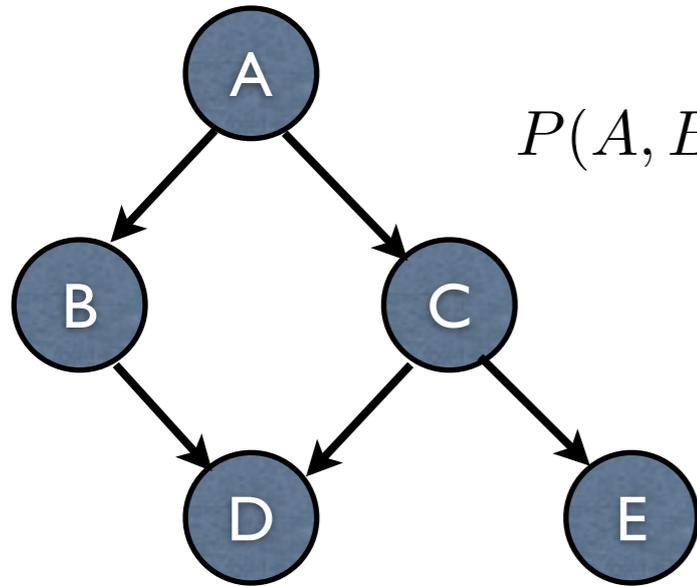
- Notes:
 - Sum over all “don’t care” variables
 - Factor common terms out of summation
 - Calculation becomes a sum of products of sums of products ...

Poly-trees are easy

- Singly-connected structures allow propagation of observations via single paths
- “Down” is just use of conditional probability
- “Up” is just Bayes rule
- Formulated as message propagation rules
- Linear time (network diameter)
- Fails on general networks!



Exact Solution of BN's (non-poly-trees)



$$P(A, B, C, D, E) = P(A)P(B|A)P(C|A)P(D|B, C)P(E|C)$$

- What is the probability of a specific state, say $A=t, B=f, C=t, D=t, E=f$?

$$p(a, \neg b, c, d, \neg e) = p(a), p(\neg b|a)p(c, a)p(d|\neg b, c)p(\neg e|c)$$

- What is the probability that $E=t$ given $B=t$?

$$p(e|b) = p(e, b)/p(b)$$

- Consider the term $P(e, b)$

$$P(e, b) = \sum_{A, C, D} P(A, b, C, D, e)$$

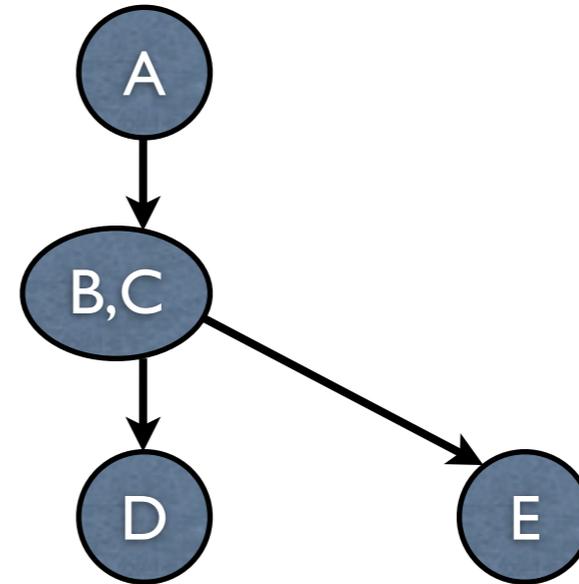
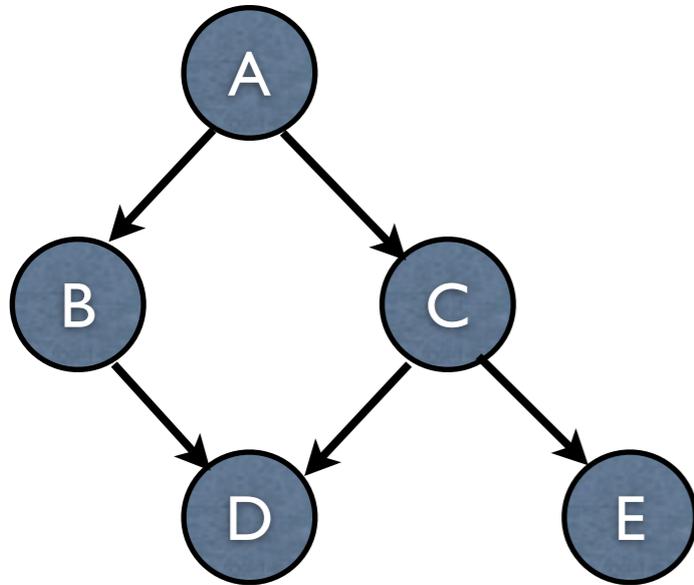
$$= \sum_{A, C, D} P(A)P(b|A)P(C|A)P(D|b, C)P(e|C)$$

$$= \sum_C P(e|C) \left(\sum_A P(A)P(C|A)P(b|A) \right) \left(\sum_D P(D|b, C) \right)$$

Alas, optimal
factoring is NP-hard

- 12 instead of 32 multiplications (even in this small example)

Other Exact Methods



- *Join-tree*: Merge variables into (small!) sets of variables to make graph into a poly-tree. Most commonly-used; aka *Clustering*, *Junction-tree*, *Potential*)
- *Cutset-conditioning*: Instantiate a (small!) set of variables, then solve each residual problem, and add solutions weighted by probabilities of the instantiated variables having those values
- ...
- All these methods are essentially equivalent; with some time-space tradeoffs.

Approximate Inference in BN's

- Direct Sampling—samples joint distribution
- Rejection Sampling—computes $P(\mathbf{X}|e)$, uses *ancestor* evidence nodes in sampling
- Likelihood Weighting—like Rejection Sampling, but weights by probability of *descendant* evidence nodes
- Markov chain Monte Carlo
 - Gibbs and other similar sampling methods

Direct Sampling

function Prior-Sample(bn) returns an event sampled from bn

inputs: bn, a Bayes net specifying the joint distribution $\mathbf{P}(X_1, \dots, X_n)$

$x :=$ an event with n elements

for $i = 1$ to n do

$x_i :=$ a random sample from $P(X_i | \text{Par}(X_i))$

return x

$$\lim_{n \rightarrow \infty} \frac{N_{PS}(x_1, \dots, x_n)}{N} = P(x_1, \dots, x_n) \quad P(x_1, \dots, x_m) \approx \frac{N_{PS}(x_1, \dots, x_m)}{N}$$

- From a large number of samples, we can estimate all joint probabilities
- The probability of an event is the fraction of all complete events generated by PS that match the partially specified event
 - hence we can compute all conditionals, etc.

Rejection Sampling

function Rejection-Sample(X , e , bn , N) returns an estimate of $P(X|e)$

inputs: bn , a Bayes net

X , the query variable

e , evidence specified as an event

N , the number of samples to be generated

local: K , a vector of counts over values of X , initially 0

for $j = 1$ to N do

$\mathbf{y} := \text{PriorSample}(bn)$

if \mathbf{y} is consistent with e then

$K[v] := K[v] + 1$ where v is the value of X in \mathbf{y}

return $\text{Normalize}(K[X])$

- Uses PriorSample to estimate the proportion of times each value of X appears in samples that are consistent with e
- But, most samples may be irrelevant to a specific query, so this is quite inefficient

Likelihood Weighting

- In trying to compute $P(X|\mathbf{e})$, where \mathbf{e} is the *evidence* (variables with known, observed values),
 - Sample only the variables other than those in \mathbf{e}
 - Weight each sample by how well it predicts \mathbf{e}

$$\begin{aligned} S_{\text{WS}}(\mathbf{z}, \mathbf{e})w(\mathbf{z}, \mathbf{e}) &= \prod_{i=1}^l P(z_i | \text{Par}(Z_i)) \prod_{i=1}^m P(e_i | \text{Par}(E_i)) \\ &= P(\mathbf{z}, \mathbf{e}) \end{aligned}$$

Likelihood Weighting

$$\begin{aligned} S_{WS}(z, e)w(z, e) &= \prod_{i=1}^l P(z_i | \text{Par}(Z_i)) \prod_{i=1}^m P(e_i | \text{Par}(E_i)) \\ &= P(z, e) \end{aligned}$$

function Likelihood-Weighting(X, e, bn, N) returns an estimate of $P(X|e)$

inputs: bn , a Bayes net

X , the query variable

e , evidence specified as an event

N , the number of samples to be generated

local: W , a vector of weighted counts over values of X , initially 0

for $j = 1$ to N do

$y, w := \text{WeightedSample}(bn, e)$

if y is consistent with e then

$W[v] := W[v] + w$ where v is the value of X in y

return $\text{Normalize}(W[X])$

function $\text{Weighted-Sample}(bn, e)$ returns an event and a weight

$x :=$ an event with n elements; $w := 1$

for $i = 1$ to n do

if X_i has a value x_i in e

then $w := w * P(X_i = x_i | \text{Par}(X_i))$

else $x_i :=$ a random sample from $P(X_i | \text{Par}(X_i))$

return x, w

Markov chain Monte Carlo

function MCMC(X, e, bn, N) returns an estimate of $P(X|e)$

local: $K[X]$, a vector of counts over values of X , initially 0

Z , the non-evidence variables in bn (includes X)

\mathbf{x} , the current state of the network, initially a copy of e

initialize \mathbf{x} with random values for the vars in Z

for $j = 1$ to N do

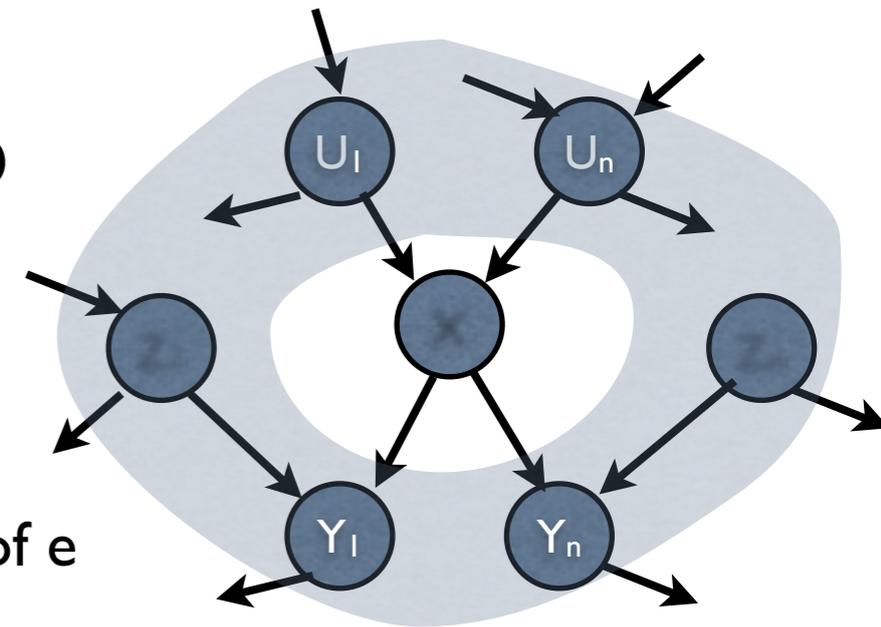
 for each Z_i in Z do

 sample the value of Z_i in \mathbf{x} from $P(Z_i|mb(Z_i))$, given the values of $mb(Z_i)$ in \mathbf{x}

$K[v] := K[v] + 1$ where v is the value of X in \mathbf{x}

return Normalize($K[X]$)

- Wander incrementally from the last state sampled, instead of re-generating a completely new sample
- For every unobserved variable, choose a new value according to its probability given the values of vars in its Markov blanket (remember, it's independent of all other vars)
- After each change, tally the sample for its value of X ; this will only change sometimes
- Problem: “narrow passages”



Most Probable Explanation

- So far, we have been solving for $P(\mathbf{X}|e)$, which yields a distribution over all possible values of the x 's
- What if we want the *best explanation* of a set of evidence, i.e., the highest-probability set of values for the x 's, given e ?
- Just *maximize* over the “don't care” variables rather than summing
- This is not necessarily the same as just choosing the value of each x with the highest probability

Rules and Probabilities

- Many have wanted to put a probability on assertions and on rules, and compute with likelihoods
- E.g., Mycin's *certainty factor* framework
 - $A (p=.3) \ \& \ B (p=.7) \implies p=.8 \implies C (p=?)$
- Problems:
 - How to combine uncertainties of preconditions and of rule
 - How to combine evidence from multiple rules
- Theorem: There is NO such algebra that works when rules are considered independently.
- Need BN for a consistent model of probabilistic inference

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