Artificial Intelligence

Lecture 7: Bayesian Network I

Xiaojin Gong 2022-04-02

Credits: AI Course in Berkeley

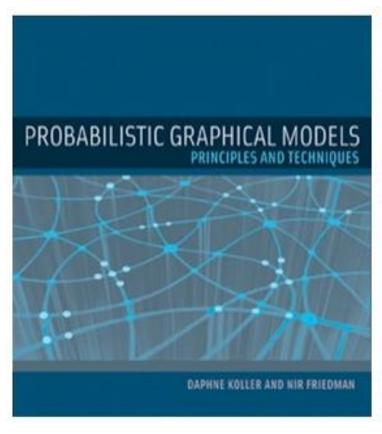
Review

- Probability
 - Random variables
 - Joint and marginal distributions
 - Conditional distribution
 - Product rule, chain rule, Bayes' rule
 - Inference
 - Independence and conditional independence

Outline

- Bayesian network
 - Representation
 - Joint probability
 - Conditional independence
 - Inference
 - Exact inference
 - Enumeration
 - Variable elimination
 - Approximate inference
 - Sampling

Reference Book





Daphne Koller Computer Science Dept. Stanford University

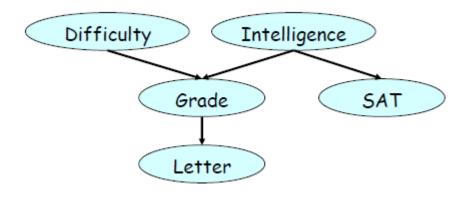


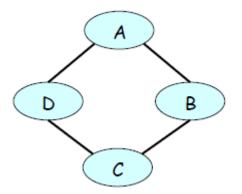
Nir Friedman
School of Computer Science &
Engineering
Hebrew University

Probabilistic Graphical Models

Bayesian network

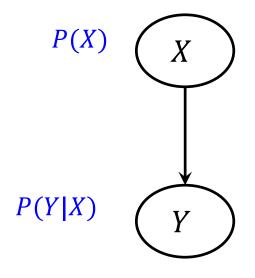






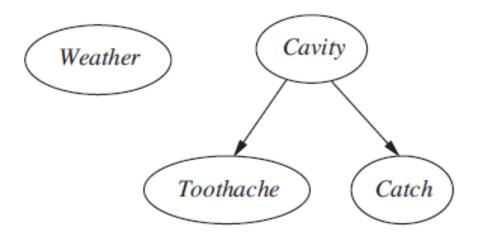
- A Bayesian network is
 - A directed acyclic graph (DAG)
 - Each node corresponds to a random variable X_i
 - Each node X_i has a conditional probability distribution (CPD) $P(X_i|Parents(X_i))$

 $A \ Bayes \ net = Topology (graph) + Local \ Conditional \ Probabilities$



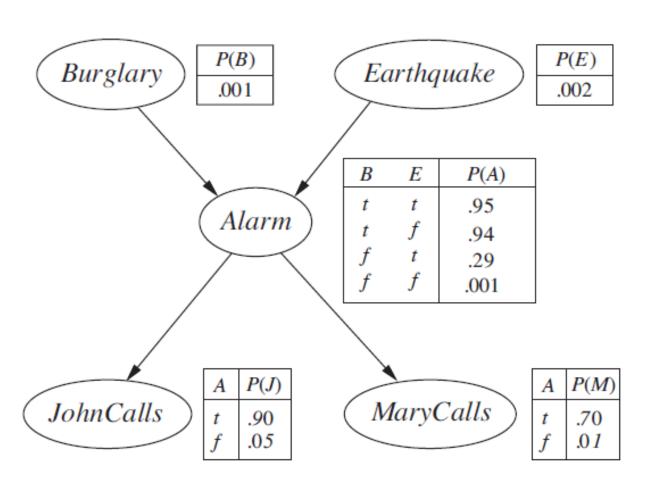
- X is a parent of Y
- X has ad direct influence on Y
- X: cause, Y:effect

• Example 1: Diagnosing a dental patient's toothache



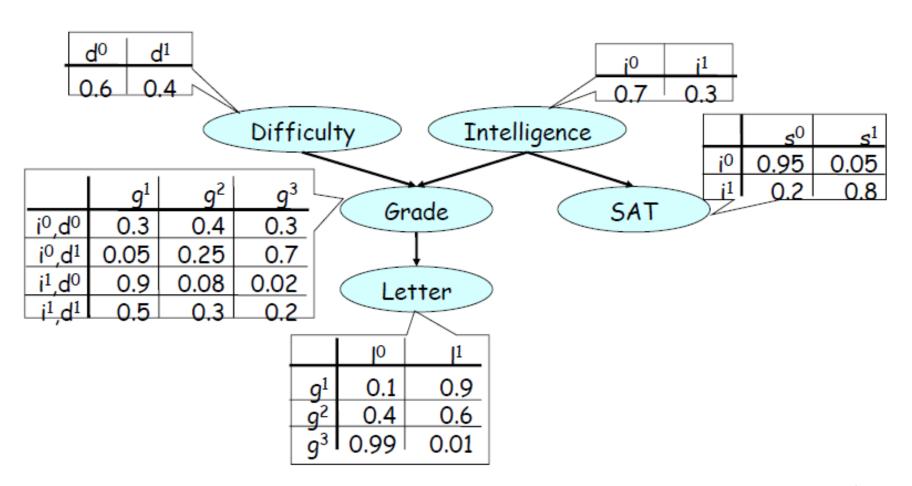
• Example 2:

Conditional probability table (CPT)

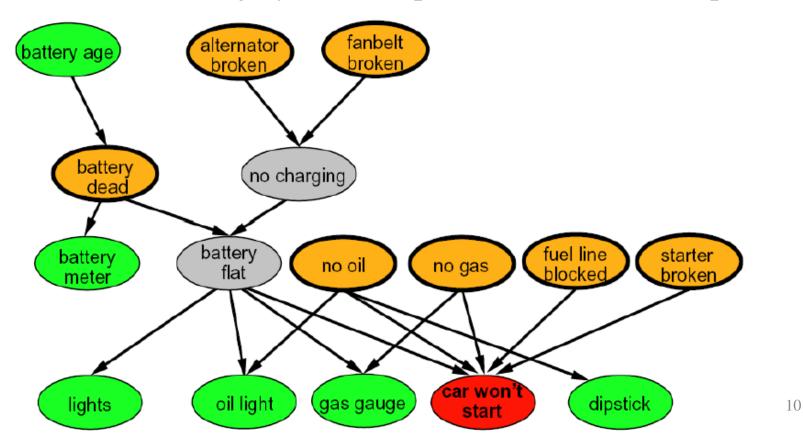


• Example 3:

Conditional probability table (CPT)



- Example 4:
 - Initial evidence: car won't start
 - Testable variables (green), "broken, so fix it" variables (orange)
 - Hidden variables (gray) ensure sparse structure, reduce parameters



Syntax:

 A BN is a directed acyclic graph with some numeric parameters attached to each node.

Semantics:

- A BN is a representation of the joint probability distribution.
- A BN is an encoding of a collection of conditional independence statements.

Joint Distribution of BN

 The BN represents a joint distribution via the chain rule for Bayesian networks

$$P(X_1, X_2, \dots, X_n) = \prod_{i} P(X_i | Parents(X_i))$$

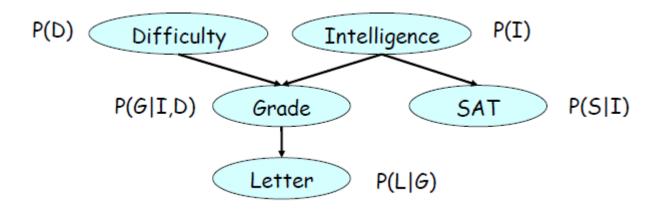
Assume conditional independences

$$P(X_i|X_1,\cdots,X_{i-1}) = P(X_i|Parents(X_i))$$

- BN is a legal distribution:
 - $P(X_1, X_2, \dots, X_n) \ge 0$

Joint Distribution of BN

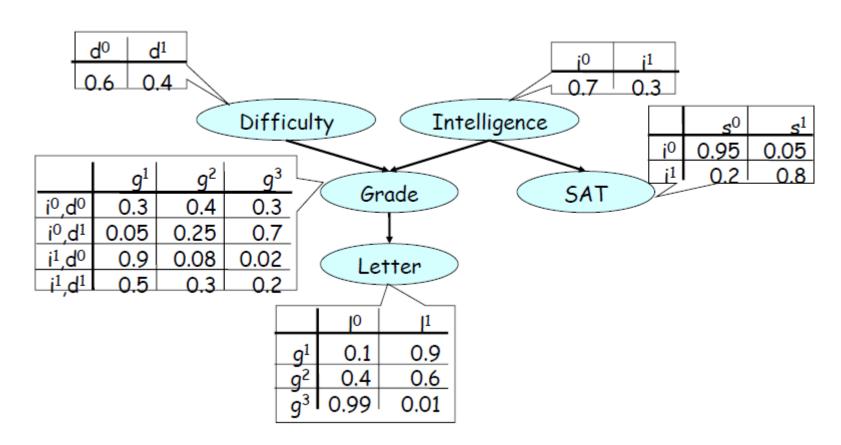
• P(D, I, G, S, L)?



$$P(D,I,G,S,L) = P(D)P(I|D)P(G|D,I)P(S|D,I,G)P(L|S,D,I,G)$$
$$= P(D)P(I)P(G|D,I)P(S|I)P(L|G)$$

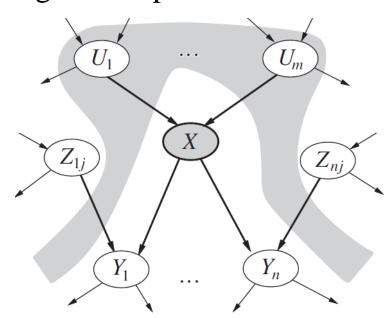
Joint Distribution of BN

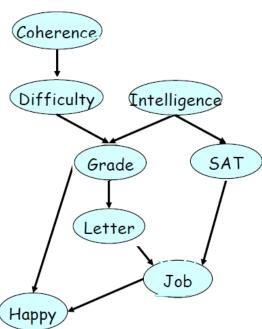
• $P(d^0, i^1, g^3, s^1, l^1)$?



- Two variables are independent $(X \perp Y)$ in a joint distribution if:
 - P(X,Y) = P(X)P(Y)
 - $\forall x, y \ P(x, y) = P(x)P(y)$
 - P(x|y) = P(x)
 - P(y|x) = P(y)
- Conditional independence: For (sets of) random variables X, Y, Z, $(X \perp Y \mid Z)$ iff:
 - P(X,Y|Z) = P(X|Z)P(Y|Z)
 - P(X|Y,Z) = P(X|Z)
 - P(Y|X,Z) = P(Y|Z)
 - $\forall x, y, z$ P(x, y|z) = P(x|z)P(y|z)

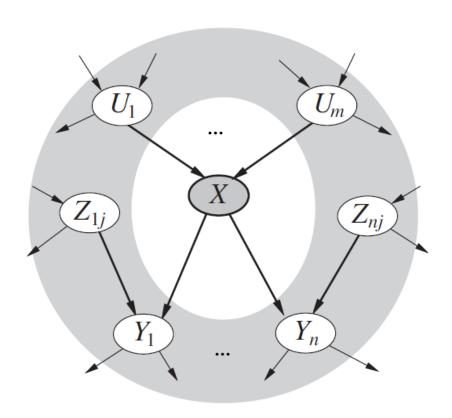
- Numerical semantics:
 - Each node is conditionally independent of its other predecessors, given its parents.
- Topological semantics:
 - Each node is conditionally independent of its non-descendants, given its parents.





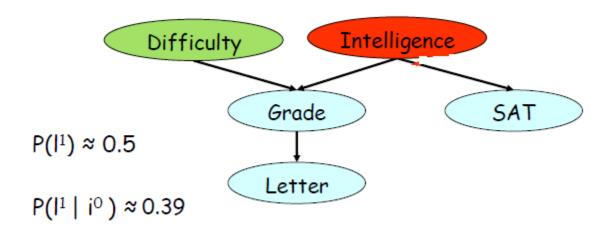
Markov blanket:

• A node is conditionally independent of all other nodes in the network, given its parents, children and children's parents

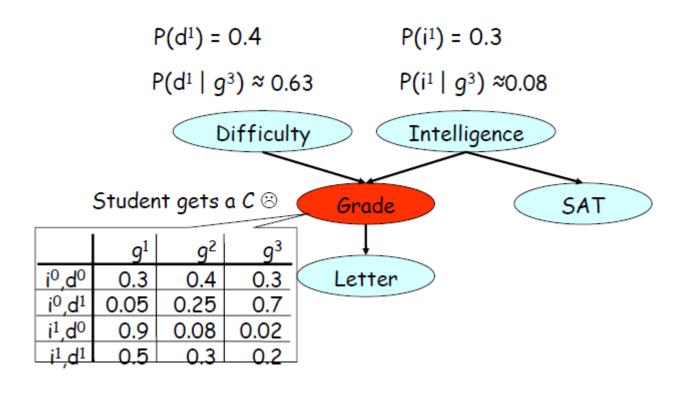


- Reasoning Patterns:
 - Important questions about a BN:
 - Are two nodes independent?
 - Are two nodes independent given certain evidence?

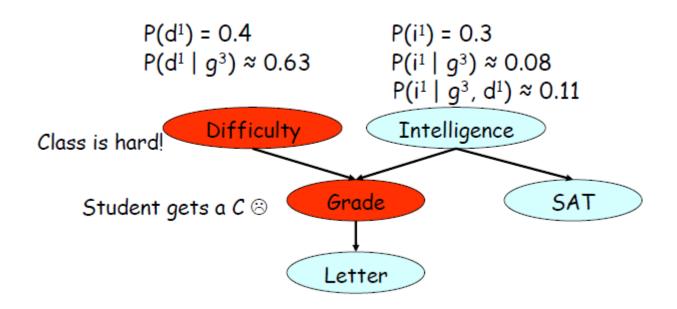
- Reasoning Patterns:
 - Causal reasoning



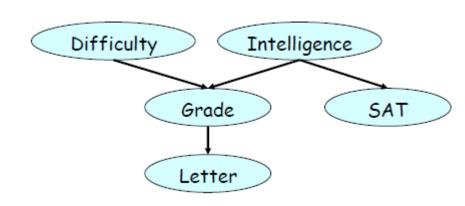
- Reasoning Patterns:
 - Evidence reasoning



- Reasoning Patterns:
 - Intercausal reasoning

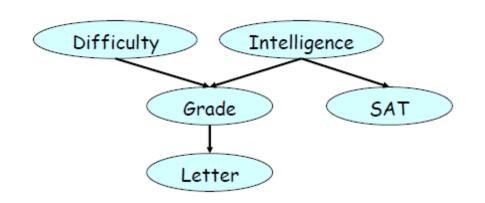


- Flow of Probabilistic Influence
 - When can X influence Y?



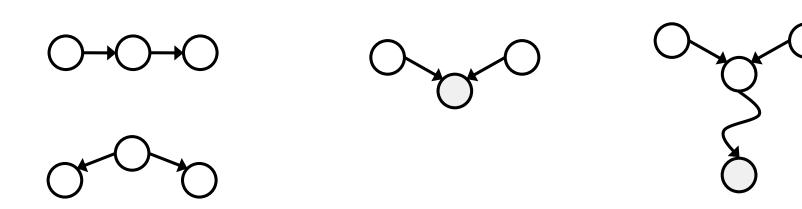
- Active trails
 - A trail $X_1 X_2 \cdots X_n$ is active if it has no v-structures $X_{i-1} \to X_i \leftarrow X_{i+1}$

- Flow of Probabilistic Influence
 - When can X influence Y given Z?
 - $X \to Y$
 - *X* ← *Y*
 - $X \to Z \to Y$
 - $X \leftarrow Z \leftarrow Y$ ×
 - $X \leftarrow Z \rightarrow Y$ ×
 - $X \to Z \leftarrow Y$

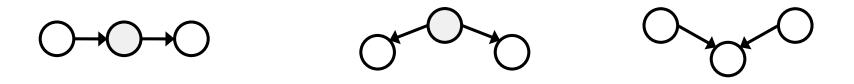


- Active trails
 - A trail $X_1 X_2 \cdots X_n$ is active given Z if
 - for any v-structures $X_{i-1} \to X_i \leftarrow X_{i+1}$ we have that X_i or one of its descendants $\in Z$
 - No other $X_i \in Z$

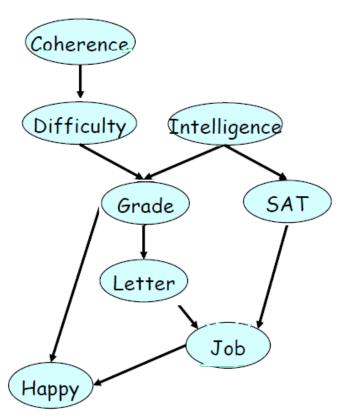
Active trails

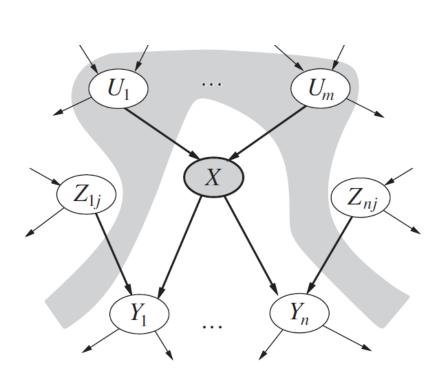


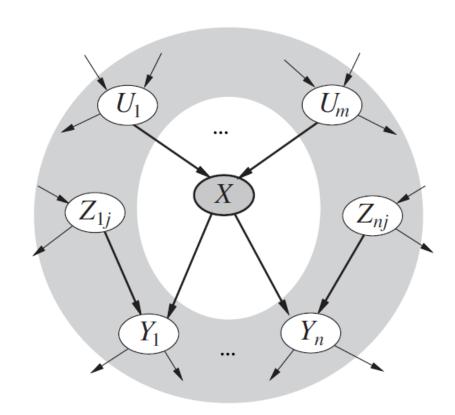
Inactive trails



- D-separation
 - $d sep_G(X_i, X_j | Z)$: X_i and X_j are d-separated in G given Z if there is no active trail in G between X and Y given Z.
 - Any node is d-separated from its non-descendants given its parents







Probabilistic Inference

- Inference:
 - Calculating some useful quantity from a joint probability distribution
- Examples:
 - Posterior probability: $P(Q|E_1 = e_1, \dots E_k = e_k)$
 - Most likely explanation: $\operatorname{argmax}_q P(Q = q | E_1 = e_1 \dots)$
- Methods:
 - Exact inference
 - Enumeration (exponential complexity)
 - Variable elimination (worst-case exponential complexity, often better)
 - Approximate inference
 - Sampling

Factors

• A factor $\Phi: (X_1, X_2, \dots, X_n) \to Val(X_1, X_2, \dots, X_n) \in R$

• Scope = $\{X_1, X_2, \dots, X_n\}$

T W P
hot sun 20
hot rain 5
cold sun 10
cold rain 15

 $\Phi(T,W)$

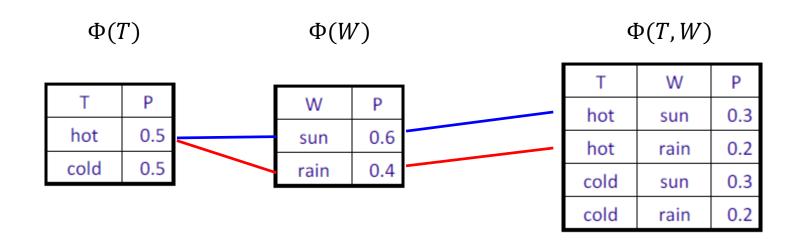
Normalize
Z = 50

Т	W	Р	
hot	sun	0.4	
hot	rain	0.1	
cold	sun	0.2	
cold	rain	0.3	

P(T,W)

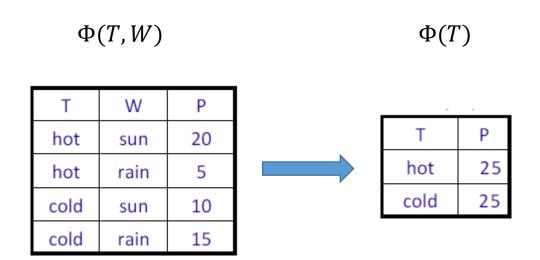
Factor Product

- $\Phi(X_1, X_2) = \Phi(X_1) \Phi(X_2)$
- $\Phi(X_1, X_2, X_3) = \Phi(X_1, X_2) \Phi(X_2, X_3)$



Factor Marginalization

•
$$\Phi(X_1) = \sum_{X_2} \Phi(X_1, X_2)$$



Enumeration

- Given
 - Evidence variables: $E_1 \dots E_k = e_1 \dots e_k$ • Query variable: Q• Hidden variables: $H_1 \dots H_r$ $X_1, X_2, \dots X_n$
- Goal: $P(Q|e_1 \dots e_k)$
- Inference by enumeration:
 - Step 1: Select the entries consistent with the evidence
 - Step 2: Sum out H to get joint of Query and evidence

$$P(Q, e_1 \dots e_k) = \sum_{h_1 \dots h_r} P(Q, h_1 \dots h_r, e_1 \dots e_k)$$

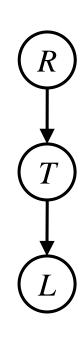
$$X_1, X_2, \dots X_n$$

Step 3: Normalization

$$P(Q|e_1\cdots e_k) = \frac{1}{Z}P(Q,e_1\cdots e_k)$$
 $Z = \sum_q P(Q,e_1\cdots e_k)$

- Random Variables
 - R: Raining
 - T: Traffic
 - L: Late for class
- Goal:

$$P(L) = ?$$



P(R)

+r	0.1
-r	0.9

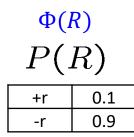
P(T|R)

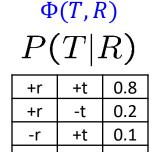
+r	+t	0.8
+r	-t	0.2
-r	+t	0.1
-r	-t	0.9

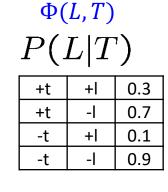
P(L|T)

+t	+	0.3
+t	- -	0.7
-t	+	0.1
-t	-	0.9

- Inference: Step 1: initialize factors & select entries
 - Initial factors are local CPTs (one per node)









- Any known values are selected
 - E.g. if we know R = +r, the initial factors are

$$P(+r)$$

$$P(T|+r)$$

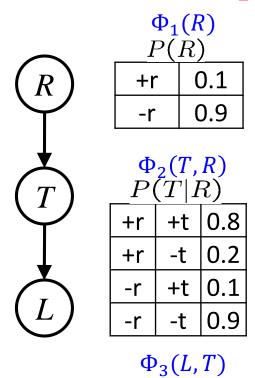
$$+r +t 0.8$$

$$+r -t 0.2$$

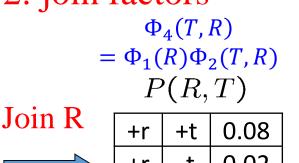
- (- -)			
+t	+	0.3	
+t	- -	0.7	
-t	+	0.1	
-t	-	0.9	

P(L|T)

• Inference: Step 2: join factors

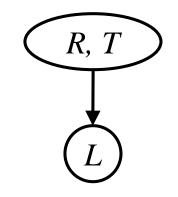


P(L T)			
+t +l 0.3			
+t	- -	0.7	
-t	+	0.1	
-t	-1	0.9	



+r	+t	0.08
+r	-t	0.02
-r	+t	0.09
-r	-t	0.81

Join T



$\Phi_5(T,R,L)$
$=\Phi_3(L,T)\Phi_4(T,R)$
P(R,T,L)

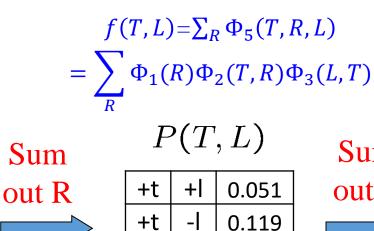
+r	+t	+	0.024
+r	+t	-	0.056
+r	-t	+1	0.002
+r	-t	-	0.018
-r	+t	+1	0.027
-r	+t	-	0.063
-r	-t	+1	0.081
-r	-t	-	0.729
			-

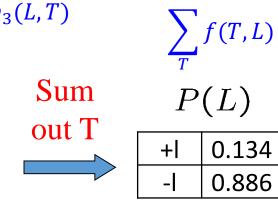
(R, T, L)

• Inference: Step 3: marginalization

$$\Phi_{5}(T,R,L)
P(R,T,L)$$

- (,,			
+r	+t	+	0.024
+r	+t	-	0.056
+r	-t	+	0.002
+r	-t	-	0.018
-r	+t	+	0.027
-r	+t	-	0.063
-r	-t	+	0.081
-r	-t	-1	0.729







0.083

0.747

Enumeration

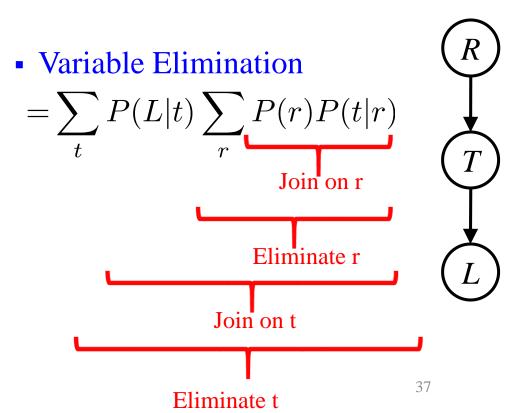
- Join up the whole joint distribution before you sum out the hidden variables
- Computational complexity: O(dⁿ)
- Very slow!

Variable Elimination

- Variable Elimination
 - Idea: interleave joining and marginalizing!
 - Still NP-hard, but usually much faster than inference by enumeration
- Inference by Enumeration

$$P(L) = \sum_{t} \sum_{r} P(L|t)P(r)P(t|r)$$
Join on t

Eliminate t

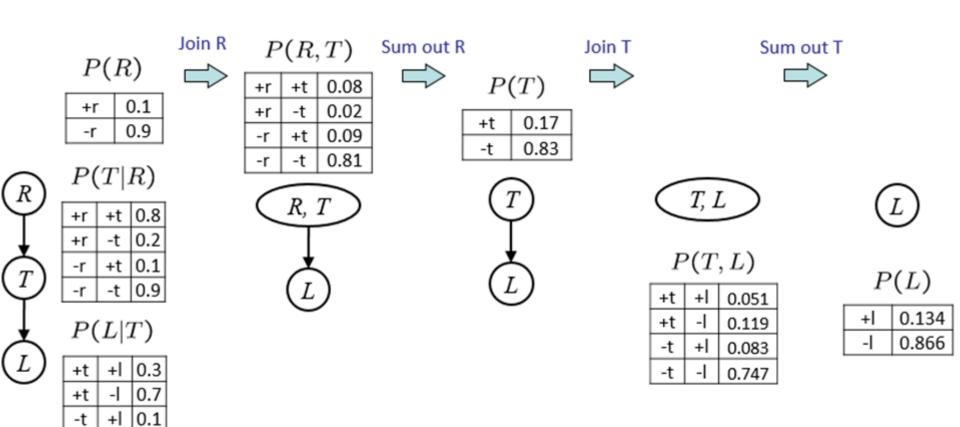


Variable Elimination

• P(L) = ?

-t

0.9



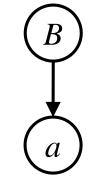
Variable Elimination with Evidence

• P(B|+a) = ?

Initial / Select



В	Р
+b	0.1
¬b	0.9





В	Α	Р
+b	+a	0.8
_		0.3
ט	⊐a	0.2
_b	+a	0.1
ک	(0.0
i D	Ia	0.5

Join on B



$$\Phi_3(B) = \Phi_1(B)\Phi_2(B)$$

$$P(a, B)$$

Α	В	Р		
+ a	+b	0.08		
+a	\negb	0.09		

Normalize

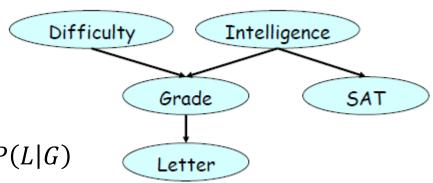


Α	В	Р
+ a	+b	8/17
+a	\negb	9/17

Variable Elimination

- Goal: P(L) = ?
- Eliminate: D, I, G, S

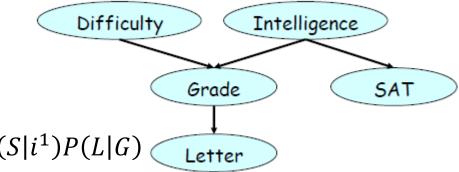




• $P(L) = \sum_{D,I,G,S} P(D,I,G,S,L)$ $= \sum_{D,I,G,S} \Phi_D(D) \Phi_I(I) \Phi_G(G,D,I) \Phi_S(S,I) \Phi_L(L,G)$ $= \sum_{I,G,S} \Phi_I(I) \Phi_S(S,I) \Phi_L(L,G) \sum_D \Phi_D(D) \Phi_G(G,D,I)$ $= \sum_{I,G,S} \Phi_I(I) \Phi_S(S,I) \Phi_L(L,G) f_1(G,I)$ $= \sum_{G,S} \Phi_L(L,G) \sum_I \Phi_I(I) \Phi_S(S,I) f_1(G,I)$ $= \sum_{G,S} \Phi_L(L,G) f_2(G,S)$ $= \sum_G \sum_S \Phi_L(L,G) f_2(G,S)$

Variable Elimination with Evidence

- Goal: $P(L, i^1) = ?$
- Eliminate: D, G, S



•
$$P(D, i^1, G, S, L) = P(D)P(i^1)P(G|D, i^1)P(S|i^1)P(L|G)$$

•
$$P(L, i^1) = \sum_{D,G,S} P(D, i^1, G, S, L)$$

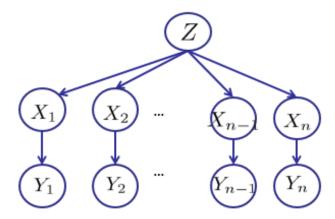
= $\sum_{D,G,S} \Phi_D(D) \Phi_G(G, D) \Phi_S(S) \Phi_L(L, G)$

Variable Elimination Algorithm

```
function ELIMINATION-ASK(X, e, bn) returns a distribution over X inputs: X, the query variable e, observed values for variables E bn, a Bayesian network specifying joint distribution P(X_1, \ldots, X_n) factors \leftarrow [] for each var in ORDER(bn.VARS) do factors \leftarrow [MAKE-FACTOR(var, e)|factors] if var is a hidden variable then factors \leftarrow SUM-OUT(var, factors) return NORMALIZE(POINTWISE-PRODUCT(factors))
```

Variable Elimination Ordering

- Query $P(X_n|y_1,...,y_n)$
- Ordering:
 - $Z, X_1, ..., X_{n-1}$ and $X_1, ..., X_{n-1}, Z$
 - What is the size of the maximum factor generated for each of the orderings?



- Answer: 2^{n+1} versus 2^2 (assuming binary)
- In general: the ordering can greatly affect efficiency.

Variable Elimination Ordering

- Computational and Space Complexity:
 - The computational and space complexity of variable elimination is determined by the largest factor
 - The elimination ordering can greatly affect the size of the largest factor.
 - Does there always exist an ordering that only results in small factors?
 - Min-neighbors
 - Min-weights

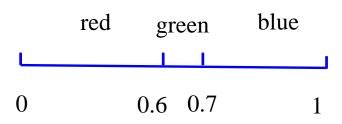
Sampling

- Basic idea
 - Draw N samples from a sampling distribution S
 - Compute an approximate posterior probability
 - Show this converges to the true probability P
- Why sampling?
 - Approximate inference: getting a sample is faster than computing the right answer (e.g. with variable elimination)

Sampling from Given Distribution

- Sampling
 - Step 1: Get sample u from uniform distribution over [0, 1)
 - Step 2: Convert this sample u into an outcome for the given distribution
- Example:
 - Discrete:

С	P(C)		
red	0.6		
green	0.1		
blue	0.3		



Sampling in Bayesian Network

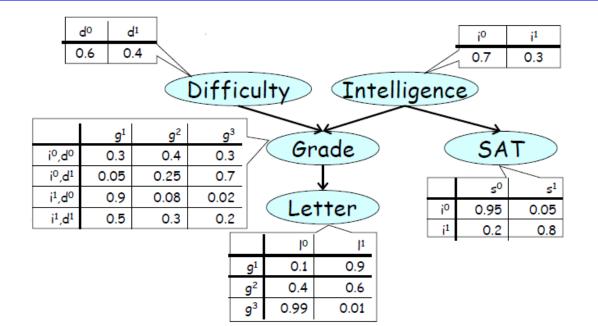
- Direct sampling methods
 - Prior / Froward sampling
 - Rejection sampling
 - Likelihood weighting
- Markov chain sampling methods
 - Gibbs sampling
 - Collapsed Gibbs sampling
 - Markov chain Monte Carlo

Prior Sampling

• Goal: Estimate P(X=x)

function PRIOR-SAMPLE(bn) returns an event sampled from the prior specified by bn inputs: bn, a Bayesian network specifying joint distribution $P(X_1, ..., X_n)$

```
\mathbf{x} \leftarrow an event with n elements foreach variable X_i in X_1, \dots, X_n do \mathbf{x}[i] \leftarrow a random sample from \mathbf{P}(X_i \mid parents(X_i)) return \mathbf{x}
```



Prior Sampling

This process generates samples with probability:

$$S_{PS}(x_1, ..., x_n) = \prod_{i=1}^{n} P(x_i | Par(X_i)) = P(x_1, ..., x_n)$$

- Let the number of samples of an event be $N_{PS}(x_1, ..., x_n)$
- Then

$$\lim_{N \to \infty} \widehat{P}(x_1, \dots, x_n) = \lim_{N \to \infty} N_{PS}(x_1, \dots, x_n) / N$$

$$= S_{PS}(x_1, \dots, x_n)$$

$$= P(x_1, \dots, x_n)$$

• The sampling procedure is consistent!

Example

• We'll get a bunch of samples from the BN:

•
$$d^0$$
, i^1 , g^1 , s^1 , l^1

•
$$d^1$$
, i^0 , g^3 , s^0 , l^0

•
$$d^0$$
, i^1 , g^2 , s^0 , l^1

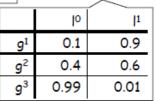
•
$$d^0$$
, i^0 , g^2 , s^0 , l^1

•
$$d^1$$
, i^1 , g^1 , s^1 , l^0

• d^0 , i^1 , g^3 , s^1 , l^0

[d ⁰	d¹					j0	j1
Ţ	0.6	0.4	1				0.7	0.3
	Difficulty Intelligence							
		g¹	g²	g ³			\rightarrow	_
i ⁰ ,d	0	0.3	0.4	0.3	Grade		SA	
i ⁰ ,d	0.0	05	0.25	0.7			7	
i ¹ ,d	0).9	0.08	0.02	Latten	-0	s ⁰	s ¹
i ¹ ,d	1 0).5	0.3	0.2	(Letter)	j ⁰	0.95	0.05
				Г	1 10 11	į1	0.2	0.8

- If we want to know P(L)
 - $l^0#:3, l^1#:3$
 - $\rightarrow P(l^0)=0.5, P(l^1)=0.5$



Rejection Sampling

• Goal: Estimate P(X=x/E=e)

```
function REJECTION-SAMPLING(X, \mathbf{e}, bn, N) returns an estimate of \mathbf{P}(X|\mathbf{e})
  inputs: X, the query variable
            e, observed values for variables E
            bn, a Bayesian network
            N, the total number of samples to be generated
  local variables: N, a vector of counts for each value of X, initially zero
  for j = 1 to N do
       \mathbf{x} \leftarrow \mathbf{PRIOR} - \mathbf{SAMPLE}(bn)
       if x is consistent with e then
         N[x] \leftarrow N[x] + 1 where x is the value of X in x
   return NORMALIZE(N)
```

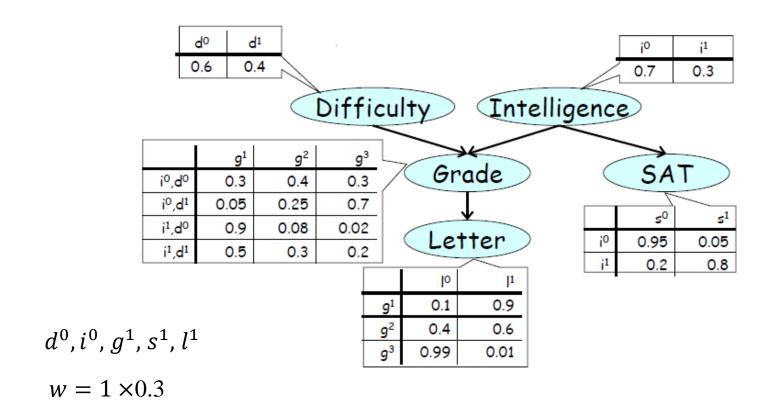
Problem: reject lots of samples if the evidence is unlikely

- Goal: Estimate P(X=x/E=e)
- Basic idea:
 - Fix evidence variables and sample the rest
 - Problem: sample distribution not consistent!
 - Solution: weight by probability of evidence given parents

• Goal: Estimate P(X=x|E=e)

```
function LIKELIHOOD-WEIGHTING(X, e, bn, N) returns an estimate of P(X|e)
  inputs: X, the query variable
            e, observed values for variables E
            bn, a Bayesian network specifying joint distribution P(X_1, \ldots, X_n)
            N, the total number of samples to be generated
  local variables: W, a vector of weighted counts for each value of X, initially zero
  for j = 1 to N do
      \mathbf{x}, w \leftarrow \mathsf{WEIGHTED\text{-}SAMPLE}(bn, \mathbf{e})
       \mathbf{W}[x] \leftarrow \mathbf{W}[x] + w where x is the value of X in x
  return NORMALIZE(W)
function WEIGHTED-SAMPLE(bn, e) returns an event and a weight
  w \leftarrow 1; \mathbf{x} \leftarrow an event with n elements initialized from e
  foreach variable X_i in X_1, \ldots, X_n do
       if X_i is an evidence variable with value x_i in e
           then w \leftarrow w \times P(X_i = x_i \mid parents(X_i))
           else \mathbf{x}[i] \leftarrow a random sample from \mathbf{P}(X_i \mid parents(X_i))
  return x, w
```

• Example: $P(L|g^1)$?



Sampling distribution if z sampled and e fixed evidence

$$S_{WS}(\mathbf{z},\mathbf{e}) = \prod_{i=1}^{l} P(z_i|\mathsf{Parents}(Z_i))$$

Now, samples have weights

$$w(\mathbf{z}, \mathbf{e}) = \prod_{i=1}^{m} P(e_i | \mathsf{Parents}(E_i))$$

Together, weighted sampling distribution is consistent

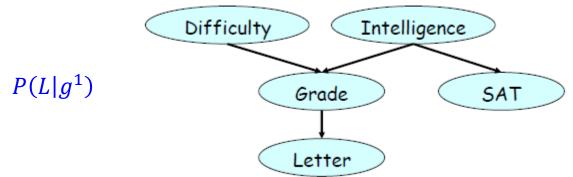
$$S_{\text{WS}}(z, e) \cdot w(z, e) = \prod_{i=1}^{l} P(z_i | \text{Parents}(z_i)) \prod_{i=1}^{m} P(e_i | \text{Parents}(e_i))$$
$$= P(\mathbf{z}, \mathbf{e})$$

Pros:

- We have taken evidence into account as we generate the sample
- More of our samples will reflect the state of the world suggested by the evidence

Cons:

- Evidence influences the choice of downstream variables, but not upstream ones
- Suffer a degradation in performance as the number of evidence variables increases.



Markov Chain Sampling

- Basic idea:
 - Generate each sample by making a random change to the preceding sample.
- Methods:
 - Gibbs sampling
 - Collapsed Gibbs sampling
 - Markov chain Monte Carlo

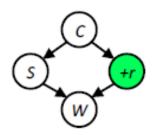
- Goal: Estimate P(X=x/E=e)
- Basic idea:
 - Step 1: Fix evidence
 - Step 2: Randomly initialize other variables
 - Step 3: Repeat
 - Choose a non-evidence variable X
 - Resample *X* from $P(X|all\ other\ variables)$

• Goal: Estimate P(X=x/E=e)

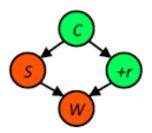
```
function GIBBS-ASK(X, \mathbf{e}, bn, N) returns an estimate of \mathbf{P}(X|\mathbf{e}) local variables: N, a vector of counts for each value of X, initially zero \mathbf{Z}, the nonevidence variables in bn \mathbf{x}, the current state of the network, initially copied from \mathbf{e} initialize \mathbf{x} with random values for the variables in \mathbf{Z} for j=1 to N do for each Z_i in \mathbf{Z} do set the value of Z_i in \mathbf{x} by sampling from \mathbf{P}(Z_i|mb(Z_i)) \mathbf{N}[x] \leftarrow \mathbf{N}[x] + 1 where x is the value of X in \mathbf{x} return NORMALIZE(N)
```

Example

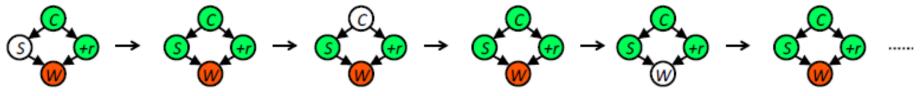
- Step 1: Fix evidence
 - R = +r



- Step 2: Initialize other variables
 - Randomly



- Steps 3: Repeat
 - Choose a non-evidence variable X
 - Resample X from P(X | all other variables)



Sample from P(S|+c,-w,+r)

Sample from P(C|+s,-w,+r)

Sample from P(W|+s,+c,+r)

- Pros:
 - The simplest Markov chain for PGMs
 - Computationally efficient to sample

- Others:
 - Collapsed Gibbs Sampling
 - Markov Chain Monte Carlo

Assignments

- Reading assignment:
 - Ch. 14.1-14.5
- Homework 4