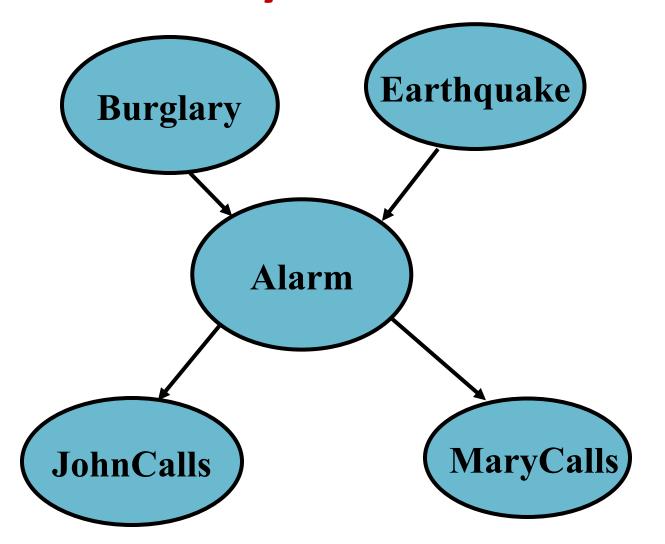
# Bayesian Networks

#### **Dr. Jianlin Cheng**

Department of Computer Science University of Missouri, Columbia

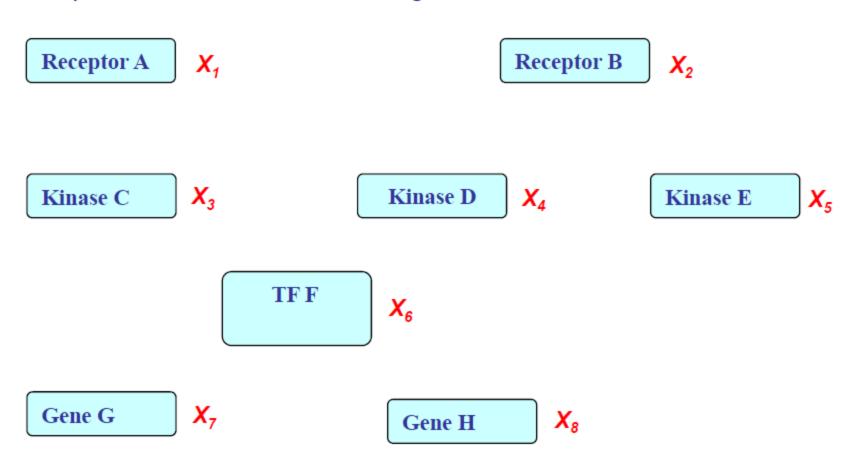
Slides Adapted from Book and CMU, MU, Stanford Machine Learning Courses
Fall, 2015

#### What is a Bayesian Network?



### What is a Bayesian Network?

A possible world for cellular signal transduction:

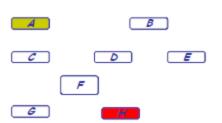


## **Basic Probability Concepts**

 Representation: what is the joint probability dist. on multiple variables?

$$P(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8)$$

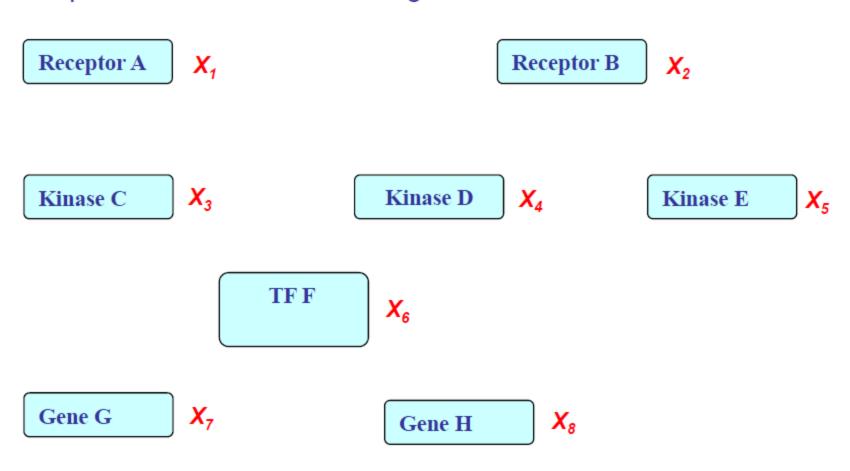
- How many state configurations in total? --- 28
- Are they all needed to be represented?
- Do we get any scientific/medical insight?



- Learning: where do we get all this probabilities?
  - Maximal-likelihood estimation? but how many data do we need?
  - Where do we put domain knowledge in terms of plausible relationships between variables, and plausible values of the probabilities?
- Inference: If not all variables are observable, how to compute the conditional distribution of latent variables given evidence?
  - Computing p(HA) would require summing over all 2<sup>6</sup> configurations of the unobserved variables

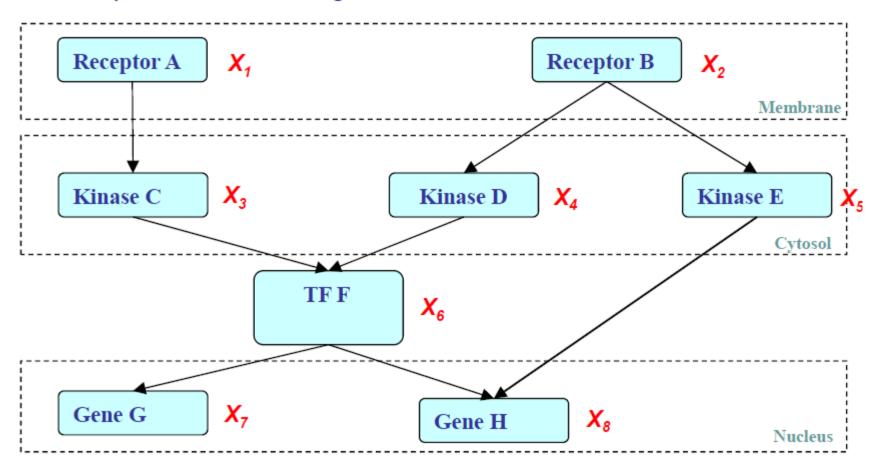
#### What is a Bayesian Network?

A possible world for cellular signal transduction:



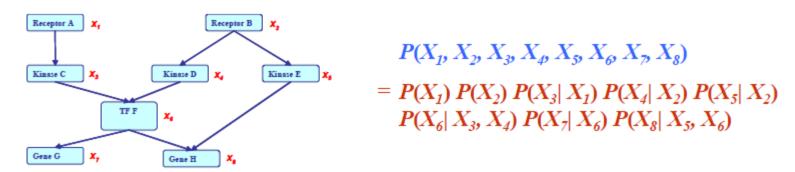
# BN: Structure Simplify Representations

Dependencies among variables



## **Bayesian Networks**

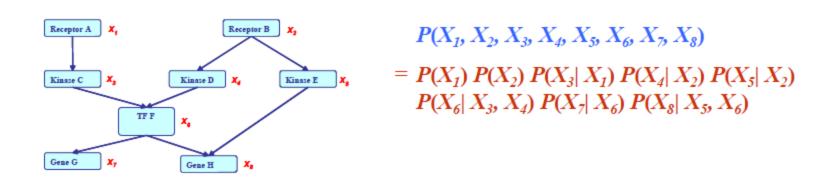
□ If  $X_i$ 's are conditionally independent (as described by a BN), the joint can be factored to a product of simpler terms, e.g.,



- Why we may favor a BN?
  - Representation cost: how many probability statements are needed?

- Algorithms for systematic and efficient inference/learning computation
  - Exploring the graph structure and probabilistic semantics
- Incorporation of domain knowledge and causal (logical) structures

# Bayesian Network: Factorization Theorem



#### Theorem:

Given a DAG, The most general form of the probability distribution that is consistent with the (probabilistic independence properties encoded in the) graph factors according to "node given its parents":

$$P(\mathbf{X}) = \prod_{i} P(X_i \mid \mathbf{X}_{\pi_i})$$

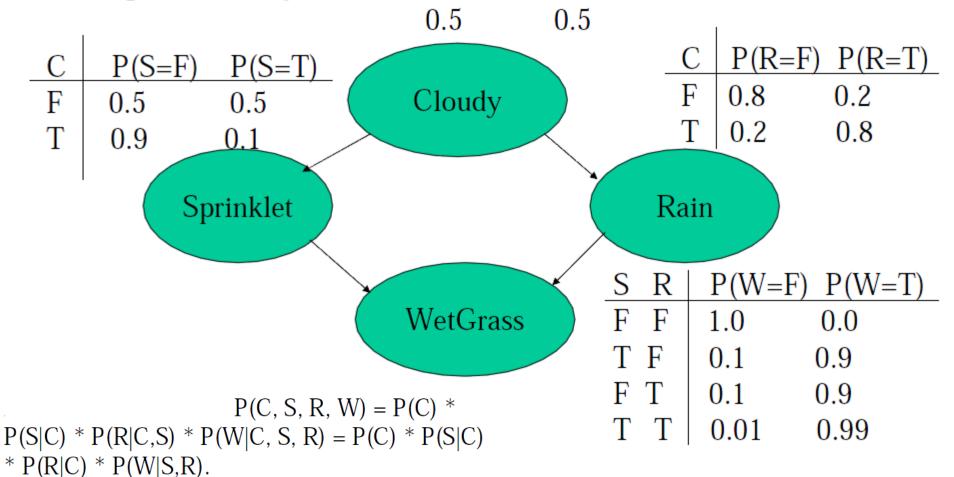
where  $X_{\pi_i}$  is the set of parents of xi. d is the number of nodes (variables) in the graph.

#### **Proof**

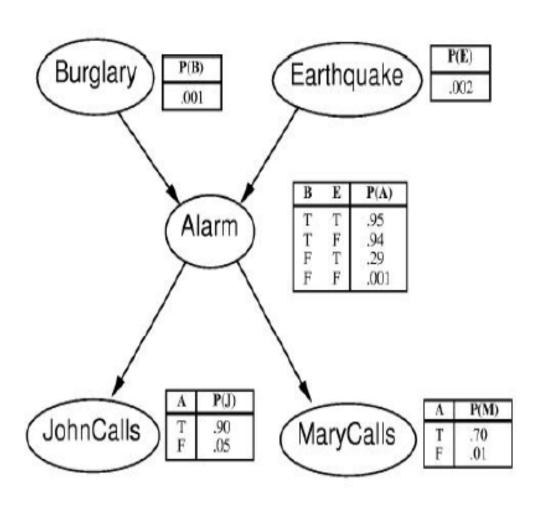
P(X<sub>1</sub>, X<sub>2</sub>, ..., X<sub>d</sub>) = P(X<sub>1</sub>|X<sub>2</sub>,X<sub>3</sub>, ..., X<sub>d</sub>) \* P(X<sub>2</sub>, X<sub>3</sub>, ..., X<sub>d</sub>) = P(X<sub>1</sub>|parent(X<sub>1</sub>)) \* P(X<sub>2</sub>|X<sub>3</sub>, ..., X<sub>d</sub>) \* P(X3, ..., Xd) = ....

#### Conditional Probability Distribution

 Discrete variable: CPT, conditional probability table <u>P(C=F)</u> <u>P(C=T)</u>



## **Examples**



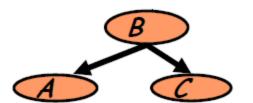
#### **Qualitative Specification**

- Where does the qualitative specification come from?
  - Prior knowledge of causal relationships
  - Prior knowledge of modular relationships
  - Assessment from experts
  - Learning from data
  - We simply link a certain architecture (e.g. a layered graph)
  - ...

#### Local Structures and Independencies

#### Common parent

Fixing B decouples A and C
 "given the level of gene B, the levels of A and C are independent"



#### Cascade

Knowing B decouples A and C
 "given the level of gene B, the level gene A provides no extra prediction value for the level of gene C"

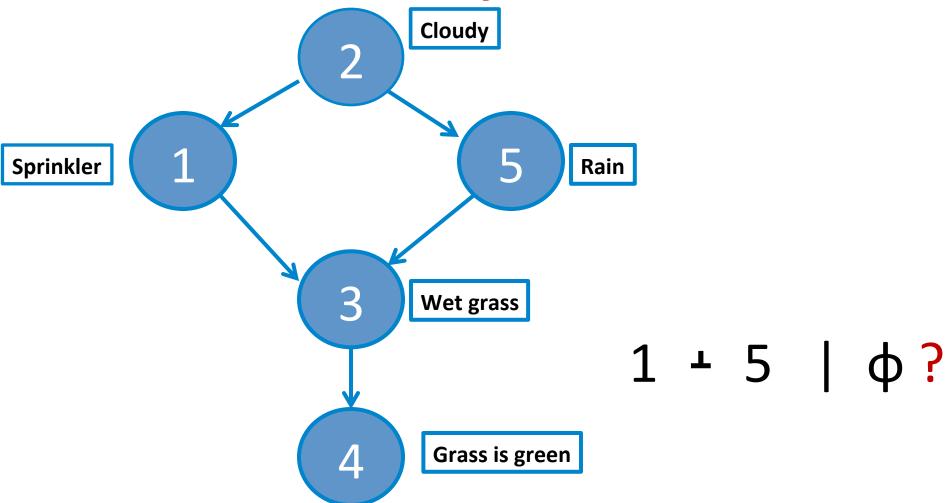


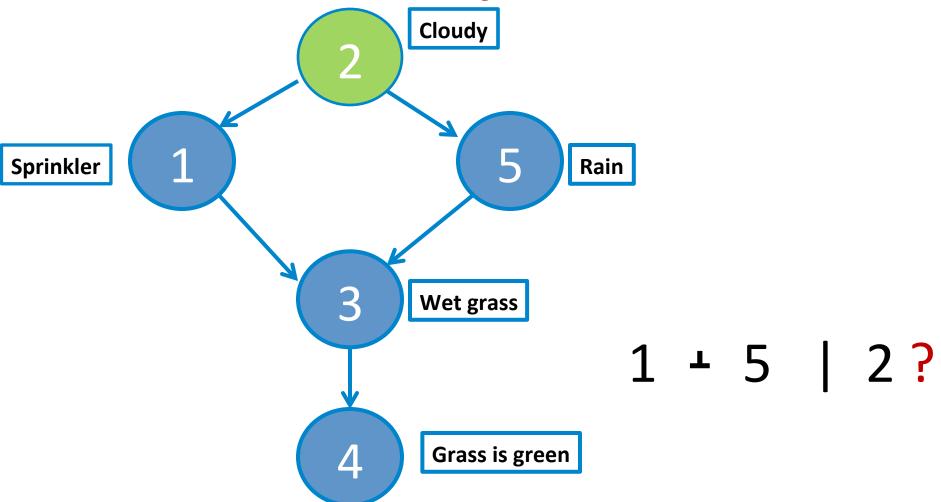
#### V-structure

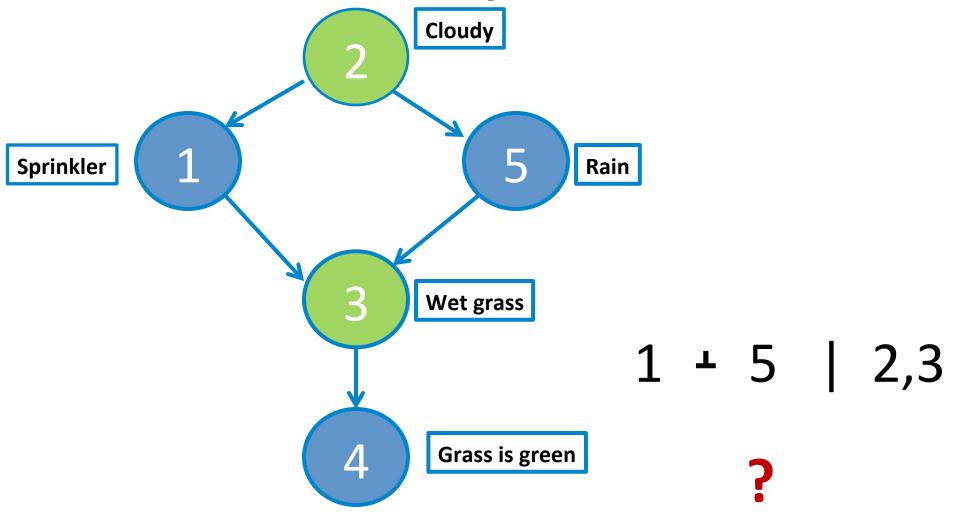
Knowing C couples A and B
 because A can "explain away" B w.r.t. C
 "If A correlates to C, then chance for B to also correlate to B will decrease"

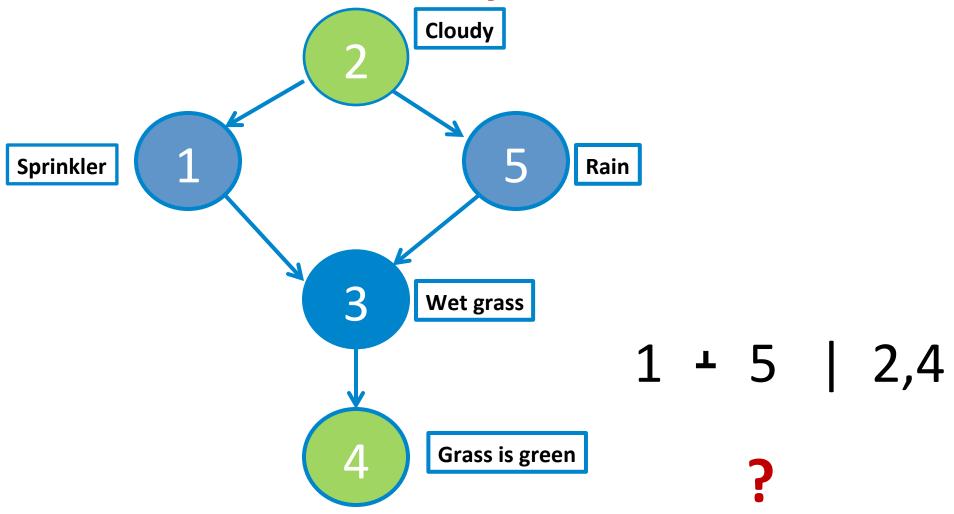


The language is compact, the concepts are rich!







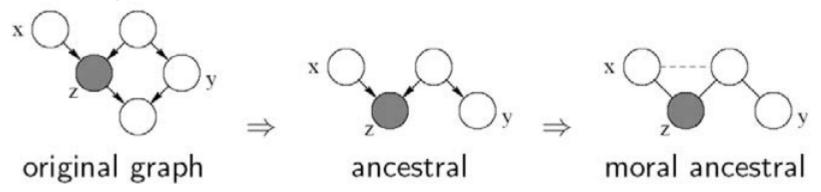


#### **Graph Separation Criterion**

 D-separation criterion for Bayesian networks (D for Directed edges):

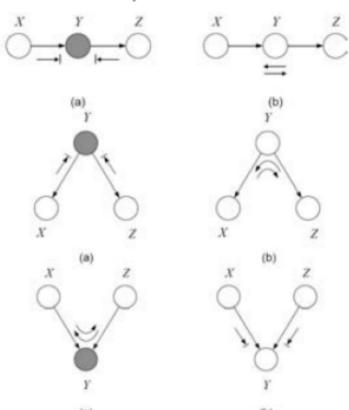
**Definition**: variables x and y are *D-separated* (conditionally independent) given z if they are separated in the *moralized* ancestral graph

Example:



## **Global Markov Properties of DAGs**

X is **d-separated** (directed-separated) from Z given Y if we can't send a ball from any node in X to any node in Z using the "*Bayes-ball*" algorithm illustrated bellow (and plus some boundary conditions):



 Defn: I(G)=all independence properties that correspond to dseparation:

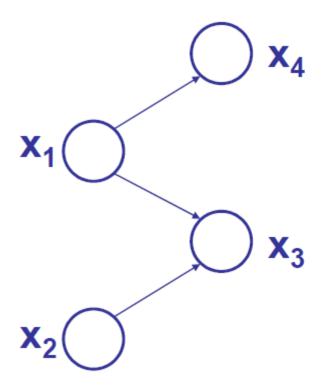
$$I(G) = \{X \perp Z | Y : dsep_G(X; Z | Y)\}$$

 D-separation is sound and complete

## **D-Separation Algorithm**

- All the paths between two nodes must be D-Separated.
- A -> B -> C (linear, B is known, then the path is blocked)
- A <- B -> C (diverging, B is known, then the path is blocked)
- A -> <u>B</u> <- C (converging, B & and its descendants are **not** known)

#### An Example



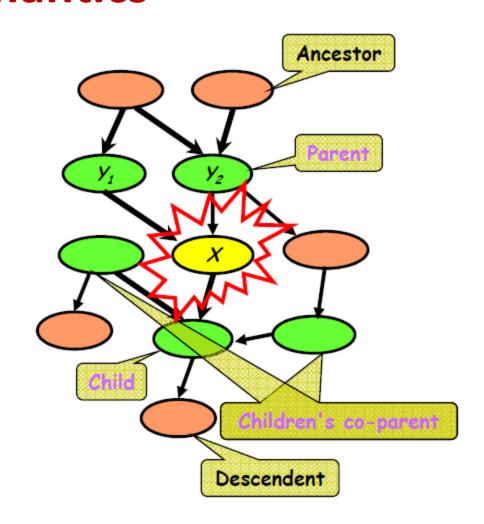
Complete the I(G) of this graph:

Essentially: A BN is a database of Pr. Independence statements among variables.

#### BN: Conditional Independence Semantics

Structure: DAG

- Meaning: a node is conditionally independent of every other node in the network outside its Markov blanket
- Local conditional distributions (CPD) and the DAG completely determine the joint dist.
- Give causality relationships, and facilitate a generative process



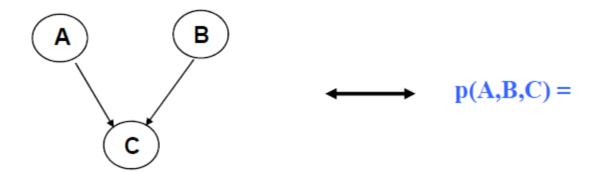
# Toward Quantitative Specification of Probability Distribution

- Separation properties in the graph imply independence properties about the associated variables
- For the graph to be useful, any conditional independence properties we can derive from the graph should hold for the probability distribution that the graph represents

#### The Equivalence Theorem

```
For a graph G,
Let \mathfrak{D}_1 denote the family of all distributions that satisfy I(G),
Let \mathfrak{D}_2 denote the family of all distributions that factor according to G,
Then \mathfrak{D}_1 \equiv \mathfrak{D}_2.
```

# **Quantitative Specification**

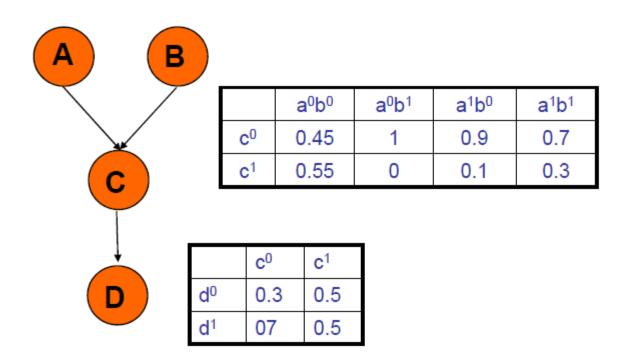


#### **Conditional Probability Tables (CPTs)**

$a^0$	0.75
a <sup>1</sup>	0.25

$b^0$	0.33
b <sup>1</sup>	0.67

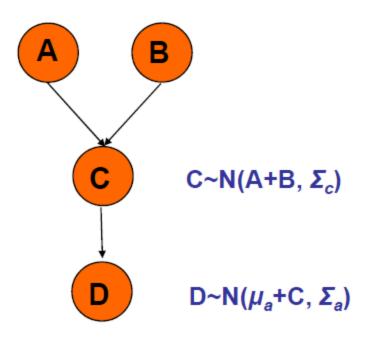
P(a,b,c.d) = P(a)P(b)P(c|a,b)P(d|c)

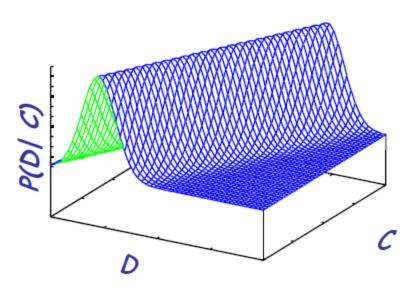


# Conditional Probability Density Function (CPDs)

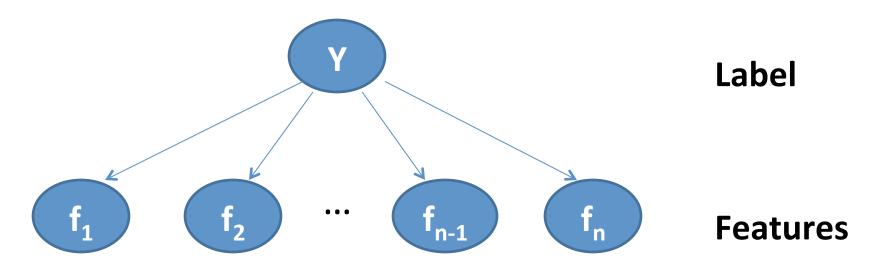
 $A \sim N(\mu_a, \Sigma_a)$   $B \sim N(\mu_b, \Sigma_b)$ 

P(a,b,c.d) = P(a)P(b)P(c|a,b)P(d|c)





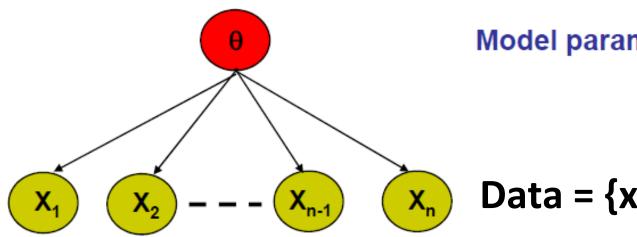
## **Conditional Independencies**



What is the model?

a)When Y is known?b)When Y is not known?

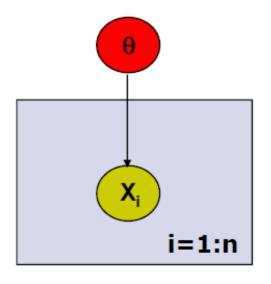
### **Conditional Independent Observations**



**Model parameters** 

Data = 
$$\{x_1, ..., X_n\}$$

#### "Plate" Notation



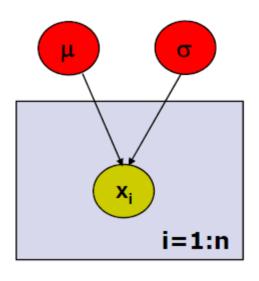
**Model parameters** 

Data = 
$$\{x_1, ..., x_n\}$$

Plate = rectangle in graphical model

variables within a plate are replicated in a conditionally independent manner

### **Example: Gaussian Model**

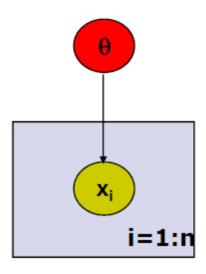


#### Generative model:

$$p(x_1,...x_n | \mu, \sigma)$$
 =  $P$   $p(x_i | \mu, \sigma)$   
=  $p(data | parameters)$   
=  $p(D | \theta)$   
where  $\theta = \{\mu, \sigma\}$ 

- Likelihood = p(data | parameters)
   = p(D | θ)
   = L (θ)
- Likelihood tells us how likely the observed data are conditioned on a particular setting of the parameters
  - Often easier to work with log L (θ)

# **Bayesian Model**



## **More Examples**

#### **Density estimation**

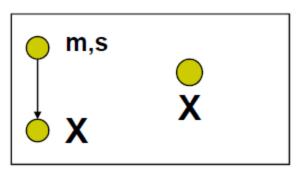
Parametric and nonparametric methods

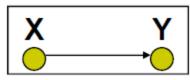
#### Regression

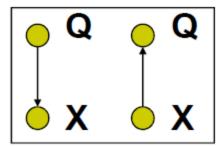
Linear, conditional mixture, nonparametric

#### Classification

Generative and discriminative approach

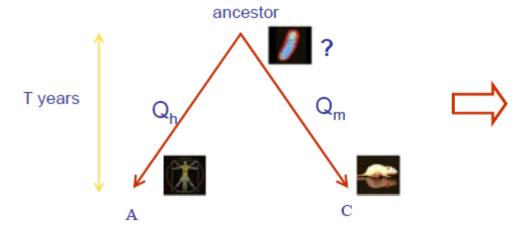


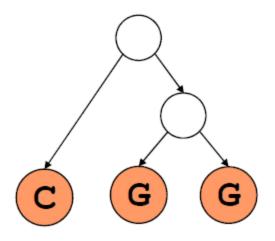




## Example, Con'd

Evolution

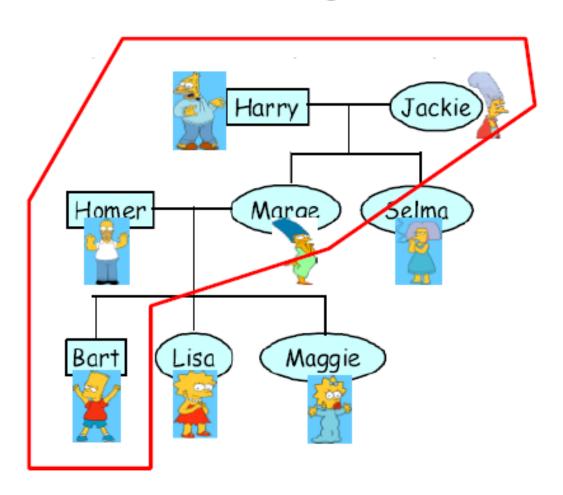




**Tree Model** 

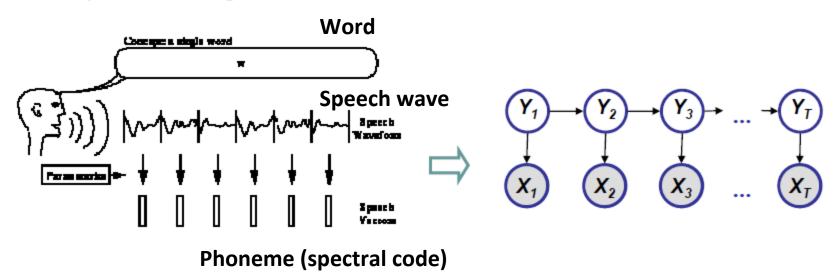
# Example, Con'd

Genetic Pedigree



# Example, Con'd

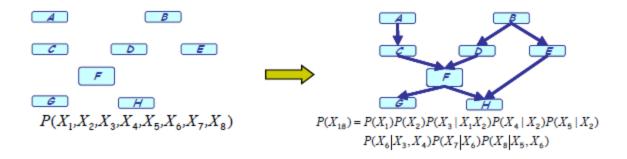
Speech recognition



**Hidden Markov Model** 

### **BN and Graphical Models**

- A Bayesian network is a special case of Graphical Models
- A Graphical Model refers to a family of distributions on a set of random variables that are compatible with all the probabilistic independence propositions encoded by a graph that connects these variables
- It is a smart way to write/specify/compose/design exponentially-large probability distributions without paying an exponential cost, and at the same time endow the distributions with structured semantics



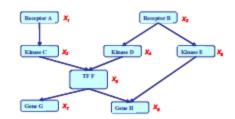
## **Two Types of GMs**

 Directed edges give causality relationships (Bayesian Network or Directed Graphical Model):

$$P(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8)$$

$$= P(X_1) P(X_2) P(X_3 | X_1) P(X_4 | X_2) P(X_5 | X_2)$$

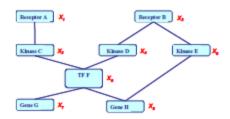
$$P(X_6 | X_3, X_4) P(X_7 | X_6) P(X_8 | X_5, X_6)$$



 Undirected edges simply give correlations between variables (Markov Random Field or Undirected Graphical model):

$$P(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8)$$

$$= \frac{1/Z} \exp\{E(X_1) + E(X_2) + E(X_3, X_1) + E(X_4, X_2) + E(X_5, X_2) + E(X_6, X_3, X_4) + E(X_7, X_6) + E(X_8, X_5, X_6)\}$$



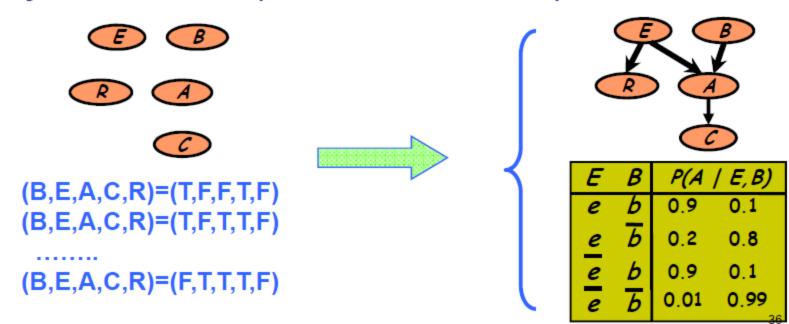
### **Probabilistic Inference**

- Computing statistical queries regarding the network, e.g.:
  - Is node X independent on node Y given nodes Z,W?
  - What is the probability of X=true if (Y=false and Z=true)?
  - What is the joint distribution of (X,Y) if Z=false?
  - What is the likelihood of some full assignment?
  - What is the most likely assignment of values to all or a subset the nodes of the network?
- General purpose algorithms exist to fully automate such computation
  - Computational cost depends on the topology of the network
  - Exact inference:
    - The junction tree algorithm
  - Approximate inference;
    - . Loopy belief propagation, variational inference, Monte Carlo sampling

## **Learning in BN**

#### The goal:

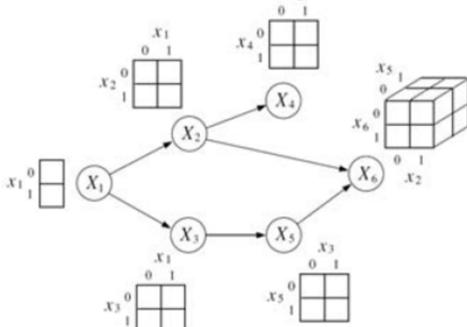
Given set of independent samples (assignments of random variables), find the best (the most likely?) Bayesian Network (both DAG and CPDs)



## **MLE Learning**

 If we assume the parameters for each CPD are globally independent, and all nodes are fully observed, then the loglikelihood function decomposes into a sum of local terms, one per node:

$$\ell(\theta; D) = \log p(D \mid \theta) = \log \prod_{w_i} \left( \prod_i p(x_{n,i} \mid \mathbf{x}_{n,\pi_i}, \theta_i) \right) = \sum_i \left( \sum_n \log p(x_{n,i} \mid \mathbf{x}_{n,\pi_i}, \theta_i) \right)$$

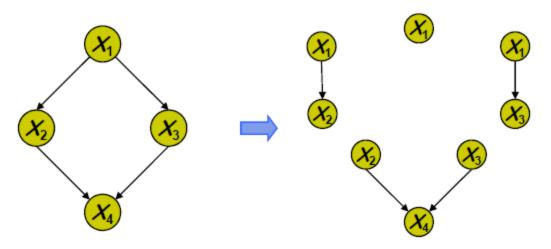


## Example: Decomposable likelihood of a directed model

Consider the distribution defined by the directed acyclic GM:

$$p(x \mid \theta) = p(x_1 \mid \theta_1) p(x_2 \mid x_1, \theta_1) p(x_3 \mid x_1, \theta_3) p(x_4 \mid x_2, x_3, \theta_1)$$

 This is exactly like learning four separate small BNs, each of which consists of a node and its parents.



### MLEs for BNs with Tabular CPDs

Assume each CPD is represented as a table (multinomial)
 where

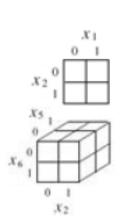
$$\theta_{ijk} \stackrel{\text{def}}{=} p(X_i = j \mid X_{\pi_i} = k)$$

- Note that in case of multiple parents, X<sub>πi</sub> will have a composite state, and the CPD will be a high-dimensional table
- The sufficient statistics are counts of family configurations

$$n_{ijk} \stackrel{\text{def}}{=} \sum\nolimits_n x_{n,i}^j x_{n,\pi_i}^k$$

- The log-likelihood is  $\ell(\theta; D) = \log \prod_{i,j,k} \theta_{ijk}^{n_{ijk}} = \sum_{i,j,k} n_{ijk} \log \theta_{ijk}$
- Using a Lagrange multiplier to enforce  $\sum_{j} \theta_{ijk} = 1$ , we get:

$$\theta_{ijk}^{ML} = \frac{n_{ijk}}{\sum_{i,j',k} n_{ij'k}}$$



## An Example

- Three variables: C Cloudy, R Rain, S –
   Sprinkler
- Data: (C=T, R = T, S = F), (C = T, R = F, S = F), (C = F, R = F, S = T)
- P(C = T) = ?, P(C = F) = ?
- $P(R = T \mid C = T) = ? P(R = F \mid C = F) = ?$
- $P(S = T \mid C = T) = ?, P(S = T \mid C = F) = ?$

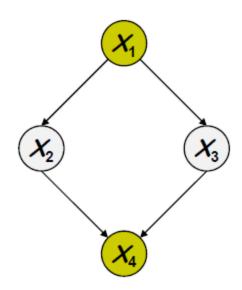
## Summary

- Represent dependency structure with a directed acyclic graph
  - Node <-> random variable
  - Edges encode dependencies
    - Absence of edge -> conditional independence
  - Plate representation
  - A BN is a database of prob. Independence statement on variables
- The factorization theorem of the joint probability
  - Local specification → globally consistent distribution
  - Local representation for exponentially complex state-space
- Support efficient inference and learning

## What if some nodes are not observed?

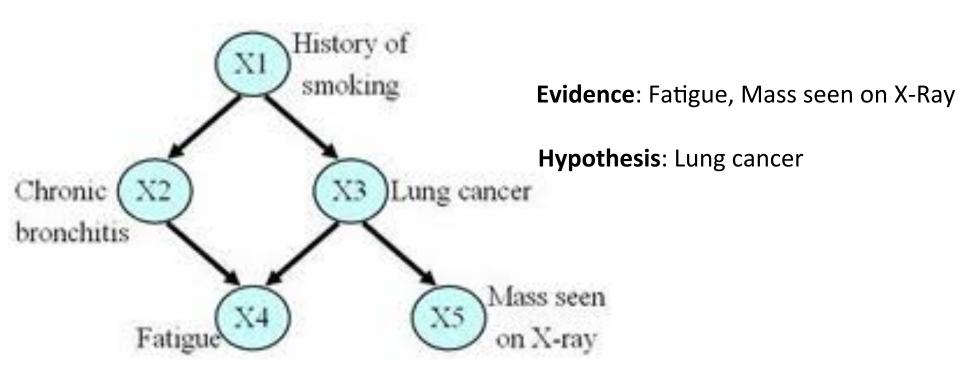
Consider the distribution defined by the directed acyclic GM:

$$p(x \mid \theta) = p(x_1 \mid \theta_1) p(x_2 \mid x_1, \theta_1) p(x_3 \mid x_1, \theta_3) p(x_4 \mid x_2, x_3, \theta_1)$$



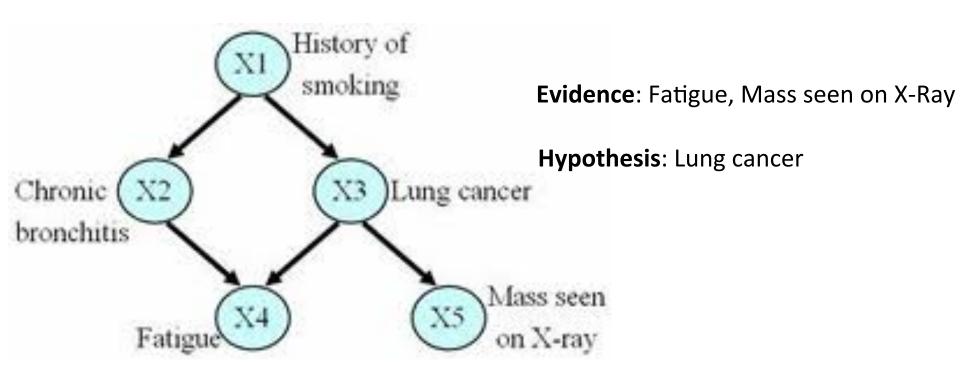
• Need to compute  $p(x_H|x_V) \rightarrow inference$ 

## **An Example**



P(Lung cancer = T | Fatigue = T, Mass X-Ray = T) = ?

## **An Example**



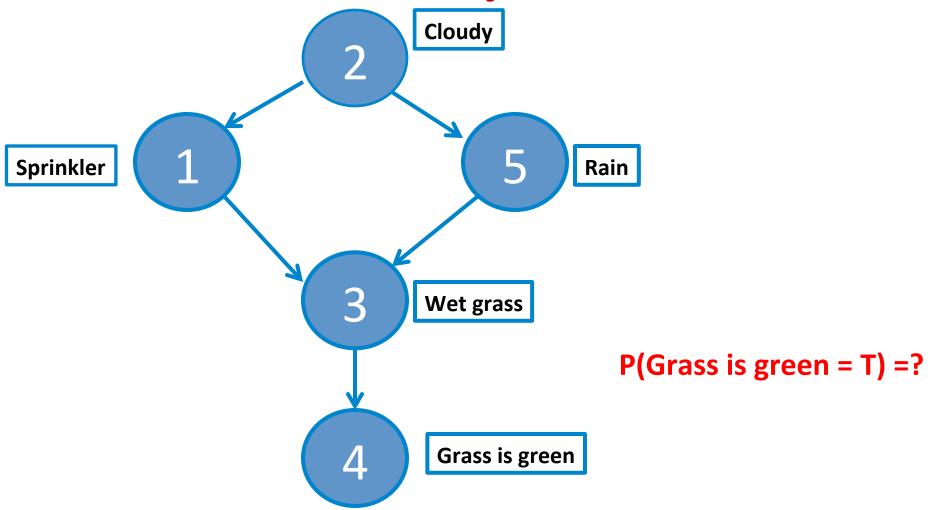
## **Inferential Query 1: Likelihood**

- Most of the queries one may ask involve evidence
  - Evidence  $\mathbf{x}_v$  is an assignment of values to a set  $\mathbf{X}_v$  of nodes in the GM over variable set  $\mathbf{X} = \{X_1, X_2, ..., X_n\}$
  - Without loss of generality X<sub>v</sub>={X<sub>k+1</sub>, ..., X<sub>n</sub>},
  - Write  $X_H = X \setminus X_v$  as the set of hidden variables,  $X_H$  can be  $\emptyset$  or X
- Simplest query: compute probability of evidence

$$P(\mathbf{X}_{\mathbf{v}}) = \sum_{\mathbf{x}_{\mathbf{H}}} P(\mathbf{X}_{\mathbf{H}}, \mathbf{X}_{\mathbf{v}}) = \sum_{x_1} \dots \sum_{x_k} P(x_1, \dots, x_k, \mathbf{X}_{\mathbf{v}})$$

this is often referred to as computing the likelihood of x<sub>v</sub>

## Assess Conditional Independence of Two Nodes in Bayesian Networks



# Inferential Query 2: Conditional Probability

 Often we are interested in the conditional probability distribution of a variable given the evidence

$$P(\mathbf{X_H} \mid \mathbf{X_V} = \mathbf{x_V}) = \frac{P(\mathbf{X_H}, \mathbf{x_V})}{P(\mathbf{x_V})} = \frac{P(\mathbf{X_H}, \mathbf{x_V})}{\sum_{\mathbf{x_H}} P(\mathbf{X_H} = \mathbf{x_H}, \mathbf{x_V})}$$

- this is the a posteriori belief in X<sub>H</sub>, given evidence x<sub>v</sub>
- We usually query a subset Y of all hidden variables X<sub>H</sub>={Y,Z} and "don't care" about the remaining, Z:

$$P(\mathbf{Y} \mid \mathbf{x}_{v}) = \sum_{\mathbf{z}} P(\mathbf{Y}, \mathbf{Z} = \mathbf{z} \mid \mathbf{x}_{v})$$

• the process of summing out the "don't care" variables z is called marginalization, and the resulting  $P(Y|X_v)$  is called a marginal prob.

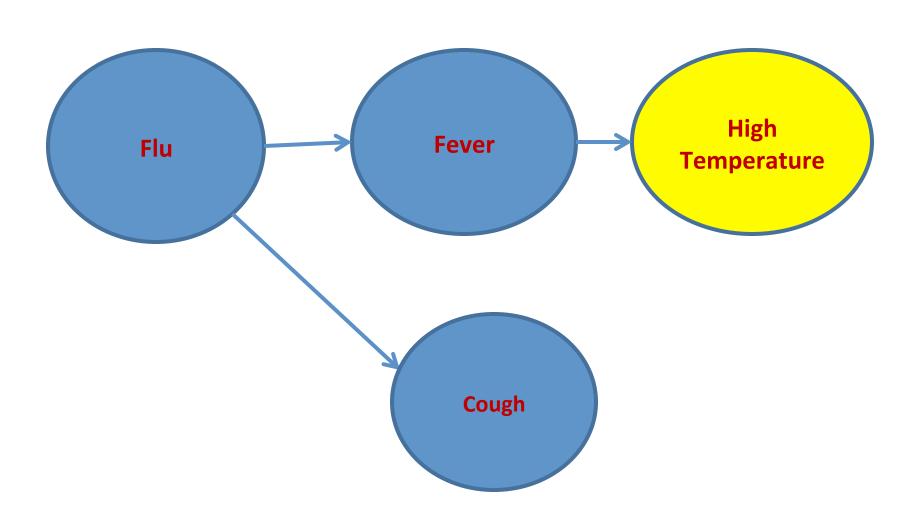
## Applications of a posterior belief

- Prediction: what is the probability of an outcome given the starting condition
  - the query node is a descendent of the evidence
- Diagnosis: what is the probability of disease/fault given symptoms

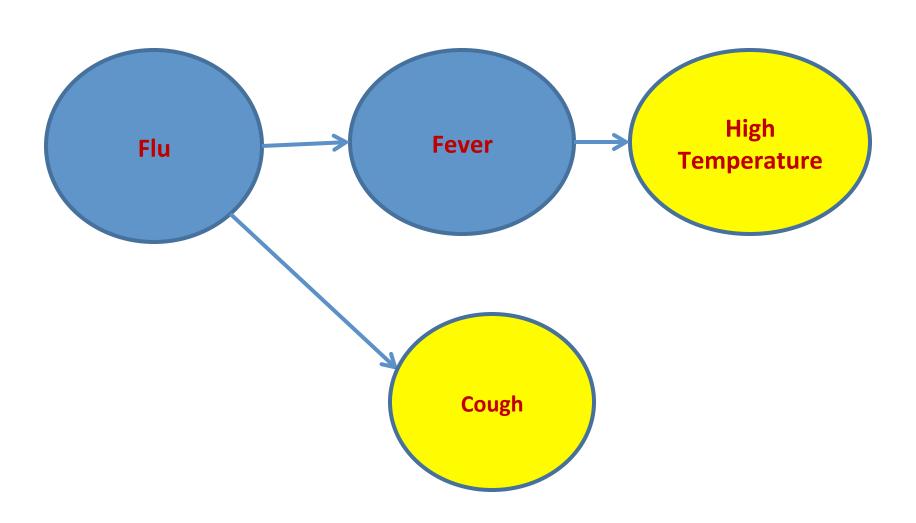


- the query node an ancestor of the evidence
- Learning under partial observation
  - fill in the unobserved values under an "EM" setting
- The directionality of information flow between variables is not restricted by the directionality of the edges in a GM
  - probabilistic inference can combine evidence form all parts of the network

## **An Example**



### **An Example – Combining Evidences**



# Inferential query 3: most probable assignment

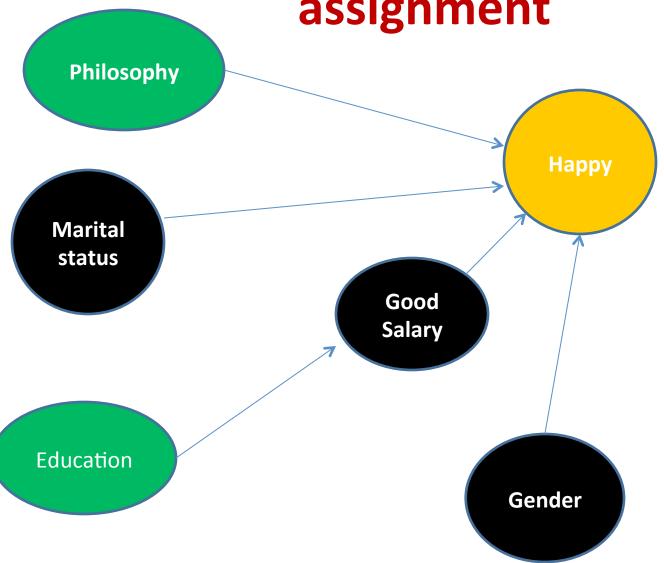
 In this query we want to find the most probable joint assignment (MPA) for some variables of interest

 Such reasoning is usually performed under some given evidence x<sub>v</sub>, and ignoring (the values of) other variables Z:

$$\mathbf{Y}^* \mid \mathbf{x}_{\mathbf{V}} = \arg\max_{\mathbf{y}} P(\mathbf{Y} \mid \mathbf{x}_{\mathbf{V}}) = \arg\max_{\mathbf{y}} \sum_{\mathbf{z}} P(\mathbf{Y}, \mathbf{Z} = \mathbf{z} \mid \mathbf{x}_{\mathbf{V}})$$

this is the maximum a posteriori configuration of Y.

# Inferential query 3: most probable assignment



## **Complexity of Inference**

#### Thm:

Computing  $P(X_H = x_H | x_v)$  in an arbitrary BN is NP-hard

- Hardness does not mean we cannot solve inference
  - It implies that we cannot find a general procedure that works efficiently for arbitrary BNs
  - For particular families of BNs, we can have provably efficient procedures

### **Approach to Inference**

- Exact inference algorithms
  - The elimination algorithm



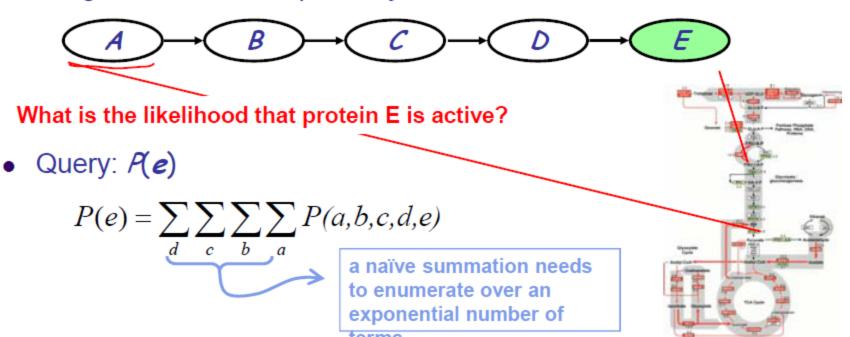
The junction tree algorithms

Approximate inference techniques

- Stochastic simulation / sampling methods
- Markov chain Monte Carlo methods
- Variational algorithms

## Marginalization and Elimination

A signal transduction pathway:



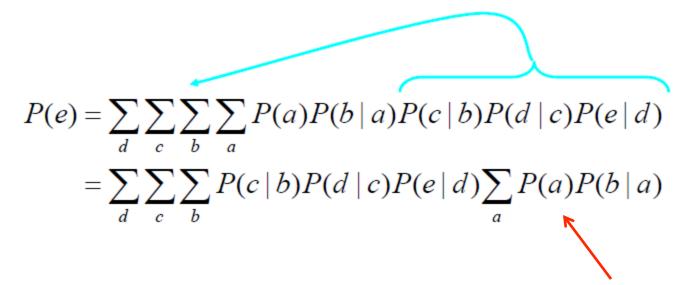
By chain decomposition, we get

$$= \sum_{d} \sum_{c} \sum_{b} \sum_{a} P(a) P(b \mid a) P(c \mid b) P(d \mid c) P(e \mid d)$$

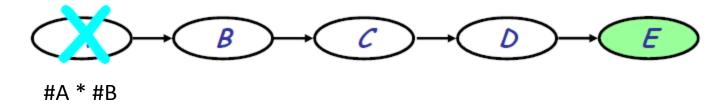
### **Elimination on Chains**



Rearranging terms ...



Only calculated once for each b, i.e. #A \* #B operations



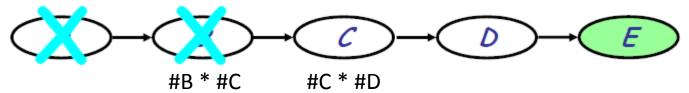
Now we can perform innermost summation

$$P(e) = \sum_{d} \sum_{c} \sum_{b} P(c | b) P(d | c) P(e | d) \sum_{a} P(a) P(b | a)$$

$$= \sum_{d} \sum_{c} \sum_{b} P(c | b) P(d | c) P(e | d) p(b)$$

 This summation "eliminates" one variable from our summation argument at a "local cost".

### **Elimination on Chains**



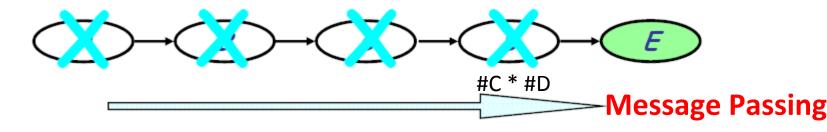
Rearranging and then summing again, we get

$$P(e) = \sum_{d} \sum_{c} \sum_{b} P(c | b) P(d | c) P(e | d) p(b)$$

$$= \sum_{d} \sum_{c} P(d | c) P(e | d) \sum_{b} P(c | b) p(b)$$

$$= \sum_{d} \sum_{c} P(d | c) P(e | d) p(c)$$

### **Elimination on Chains**



Eliminate nodes one by one all the way to the end, we get

$$P(e) = \sum_{d} P(e \mid d) p(d)$$

- Complexity:
  - Each step costs  $O(|Val(X_i)|^*|Val(X_{i+1})|)$  operations:  $O(nk^2)$
  - Compare to naïve evaluation that sums over joint values of n-1 variables  $O(k^n)$

## Inference on General BN via Variable Elimination

#### General idea:

Write query in the form

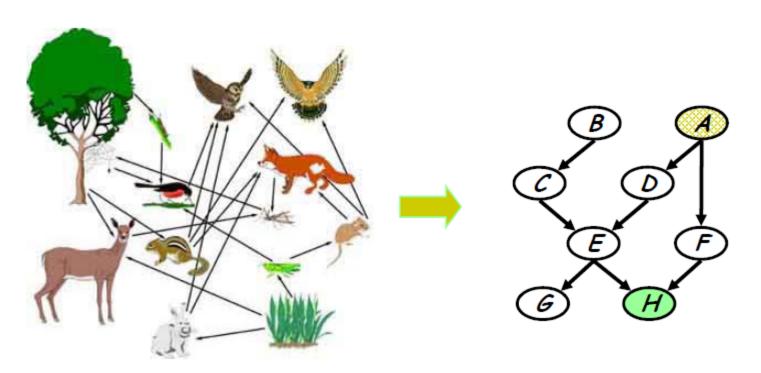
$$P(X_1, \mathbf{e}) = \sum_{x_n} \cdots \sum_{x_3} \sum_{x_2} \prod_i P(x_i \mid pa_i)$$

- this suggests an "elimination order" of latent variables to be marginalized
- Iteratively
  - Move all irrelevant terms outside of innermost sum
  - Perform innermost sum, getting a new term
  - Insert the new term into the product
- wrap-up

$$P(X_1 | e) = \frac{P(X_1, e)}{P(e)}$$

## A more complex network

#### A food web

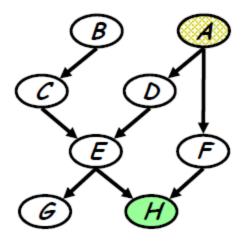


What is the probability that hawks are leaving given that the grass condition is poor?

- Query: P(A | h)
  - Need to eliminate: B,C,D,E,F,G,H
- Initial factors:

$$P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)P(g \mid e)P(h \mid e,f)$$

Choose an elimination order: H,G,F,E,D,C,B

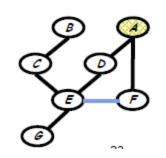


- Step 1:
  - Conditioning (fix the evidence node (i.e., h) on its observed value (i.e., h):

$$m_h(e, f) = p(h = \widetilde{h} \mid e, f)$$

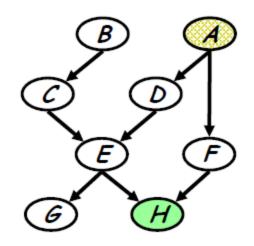
This step is isomorphic to a marginalization step:

$$m_h(e, f) = \sum_h p(h \mid e, f) \delta(h = \widetilde{h})$$



- Query: P(B | h)
  - Need to eliminate: B,C,D,E,F,G
- Initial factors:

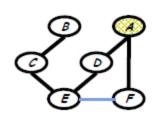
$$P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)P(g \mid e)P(h \mid e,f)$$
  
$$\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)P(g \mid e)m_h(e,f)$$



- Step 2: Eliminate G
  - compute

$$m_g(e) = \sum_g p(g \mid e) = 1$$

- $\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)m_g(e)m_h(e,f)$
- $= P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)m_h(e, f)$

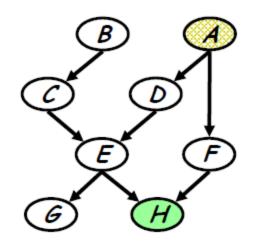


- Query: P(B | h)
  - Need to eliminate: B,C,D,E,F
- Initial factors:

$$P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)P(g \mid e)P(h \mid e,f)$$

$$\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)P(g \mid e)m_h(e,f)$$

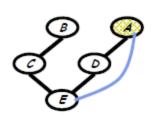
$$\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)P(f \mid a)m_h(e,f)$$



- Step 3: Eliminate F
  - compute

$$m_f(e,a) = \sum_f p(f \mid a) m_h(e,f)$$

 $\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c,d)m_f(a,e)$ 



Calculations: #F \* (#E \* #A)

- Query: P(B | h)
  - Need to eliminate: B,C,D,E
- Initial factors:

$$P(a)P(b)P(c\mid b)P(d\mid a)P(e\mid c,d)P(f\mid a)P(g\mid e)P(h\mid e,f)$$

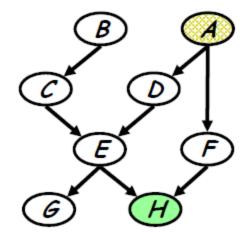
- $\Rightarrow P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)m_h(e,f)$
- $\Rightarrow P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)m_h(e,f)$
- $\Rightarrow P(a)P(b)P(c|b)P(d|a)P(e|c,d)m_f(a,e)$

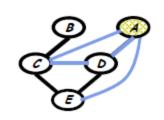


compute

$$m_e(a,c,d) = \sum_e p(e \mid c,d) m_f(a,e)$$

 $\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)m_e(a,c,d)$ 



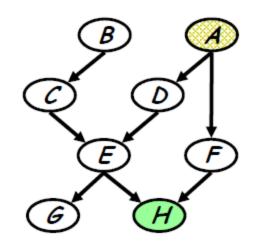


Calculations: #E \* (#A \* #C \* #D)

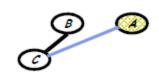
- Query: P(B | h)
  - Need to eliminate: B,C,D
- Initial factors:

$$P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f)$$

- $\Rightarrow P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)m_h(e,f)$
- $\Rightarrow P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)m_h(e,f)$
- $\Rightarrow P(a)P(b)P(c \mid b)P(d \mid a)P(e \mid c, d)m_f(a, e)$
- $\Rightarrow P(a)P(b)P(c|b)P(d|a)m_e(a,c,d)$



- Step 5: Eliminate D
  - compute  $m_d(a,c) = \sum_d p(d \mid a) m_e(a,c,d)$



 $\Rightarrow P(a)P(b)P(c \mid d)m_d(a,c)$ 

Calculations: #D \* (#A \* #C)

- Query: P(B | h)
  - Need to eliminate: B,C
- Initial factors:

$$P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)P(g \mid e)P(h \mid e, f)$$

- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)P(g \mid e)m_h(e, f)$
- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c,d)P(f \mid a)m_h(e,f)$
- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c,d)m_f(a,e)$
- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)m_e(a,c,d)$
- $\Rightarrow P(a)P(b)P(c \mid d)m_d(a,c)$

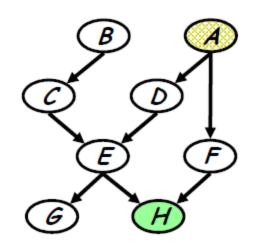




$$m_c(a,b) = \sum p(c \mid b) m_d(a,c)$$

$$\Rightarrow P(a)P(b)P(c \mid d)m_d(a,c)$$

Calculations: #C \* (#A \* #B)





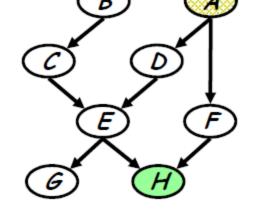
- Query: P(B | h)
  - Need to eliminate: B
- Initial factors:

$$P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)P(g \mid e)P(h \mid e, f)$$

- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)P(g \mid e)m_h(e, f)$
- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)m_h(e, f)$
- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c,d)m_f(a,e)$
- $\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)m_{s}(a,c,d)$
- $\Rightarrow P(a)P(b)P(c \mid d)m_d(a,c)$
- $\Rightarrow P(a)P(b)m_c(a,b)$
- Step 7: Eliminate B
  - compute

$$m_b(a) = \sum_b p(b) m_c(a,b)$$

 $\Rightarrow P(a)m_b(a)$ 





Calculations: #B \* #A

- Query: P(B | h)
  - Need to eliminate: B
- Initial factors:

$$P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c,d)P(f \mid a)P(g \mid e)P(h \mid e,f)$$

$$\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)P(g \mid e)m_h(e, f)$$

$$\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)P(f \mid a)m_h(e, f)$$

$$\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)P(e \mid c, d)m_f(a, e)$$

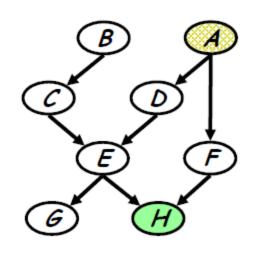
$$\Rightarrow P(a)P(b)P(c \mid d)P(d \mid a)m_{e}(a,c,d)$$

$$\Rightarrow P(a)P(b)P(c \mid d)m_d(a,c)$$

$$\Rightarrow P(a)P(b)m_c(a,b)$$

$$\Rightarrow P(a)m_b(a)$$

Step 8: Wrap-up 
$$p(a,\widetilde{h}) = p(a)m_b(a), \quad p(\widetilde{h}) = \sum_a p(a)m_b(a)$$
$$\Rightarrow P(a \mid \widetilde{h}) = \frac{p(a)m_b(a)}{\sum_b p(a)m_b(a)}$$



# Complexity of Variable Elimination

Suppose in one elimination step we compute

$$m_x(y_1,...,y_k) = \sum_x m'_x(x,y_1,...,y_k)$$
  
 $m'_x(x,y_1,...,y_k) = \prod_{i=1}^k m_i(x,\mathbf{y}_{c_i})$ 

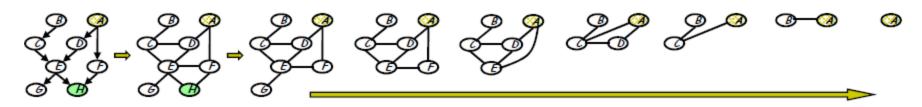
#### This requires

- $k \bullet |Val(X)| \bullet \prod_{i} |Val(\mathbf{Y}_{C_i})|$  multiplications
  - For each value of  $x_1, y_2, ..., y_k$ , we do k multiplications
- $|Val(X)| \bullet \prod_{i} |Val(\mathbf{Y}_{C_i})|$  additions
  - For each value of  $y_1$ , ...,  $y_k$ , we do |Val(X)| additions

Complexity is **exponential** in number of variables in the intermediate factor

### **Understanding Variable Elimination**

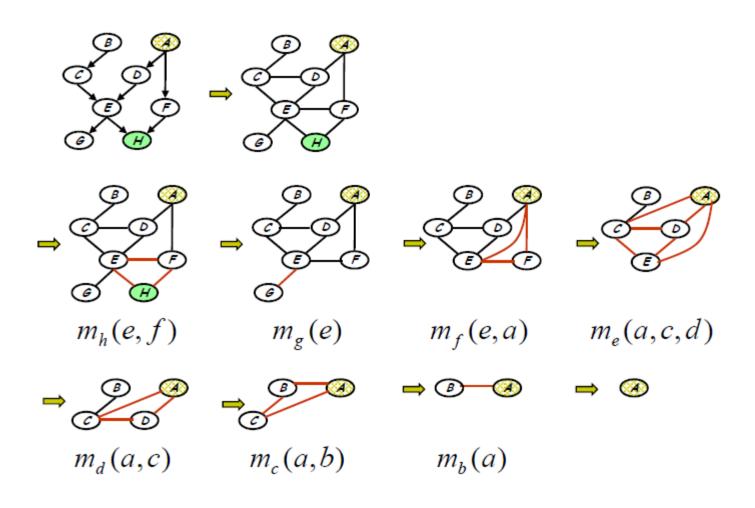
A graph elimination algorithm



moralization

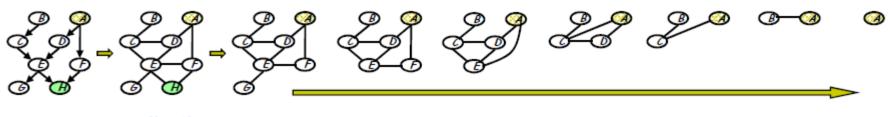
graph elimination

## **Elimination Cliques**



### **Understanding Variable Elimination**

A graph elimination algorithm

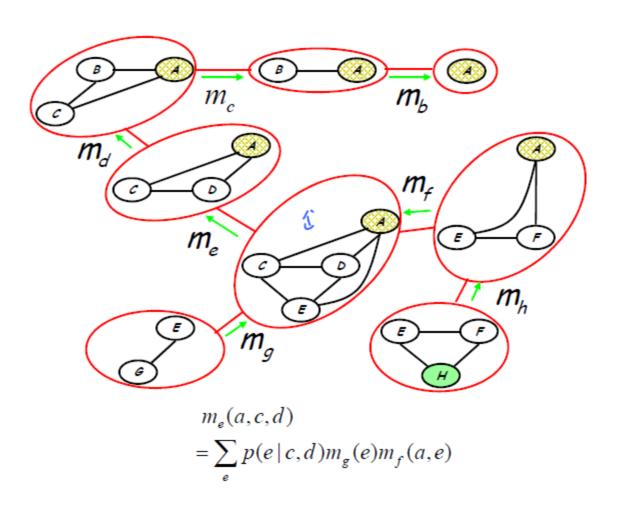


moralization

graph elimination

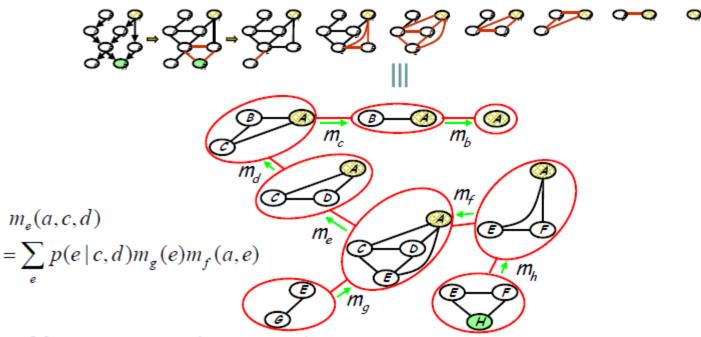
- Intermediate terms correspond to the cliques resulted from elimination
  - "good" elimination orderings lead to small cliques and hence reduce complexity (what will happen if we eliminate "e" first in the above graph?)
  - finding the optimum ordering is NP-hard, but for many graph optimum or nearoptimum can often be heuristically found
- Applies to undirected GMs

## **A Clique Tree**



### From Elimination to Message Passing

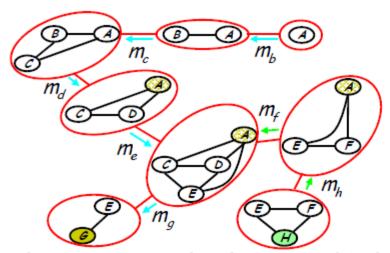
- Our algorithm so far answers only one query (e.g., on one node), do we need to do a complete elimination for every such query?
- Elimination = message passing on a clique tree



Messages can be reused

### From Elimination to Message Passing

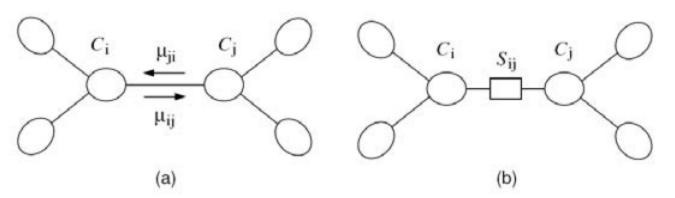
- Our algorithm so far answers only one query (e.g., on one node), do we need to do a complete elimination for every such query?
- Elimination = message passing on a clique tree
  - Another query ...



• Messages  $m_f$  and  $m_h$  are reused, others need to be recomputed

## **The Junction Tree Algorithm**

Shafer-Shenoy algorithm



Message from clique / to clique j :

Potential of C<sub>i</sub> itself

$$\mu_{i \to j} = \sum_{C_i \setminus S_{ii}} \psi_{C_i} \prod_{k \neq j} \mu_{k \to i}(S_{ki})$$

Clique marginal

$$p(C_i) \propto \psi_{C_i} \prod_k \mu_{k \to i}(S_{ki})$$

Message passed Into i from all sources Except j

**Probability of C<sub>i</sub> = its potential \* messages coming from all sources** 

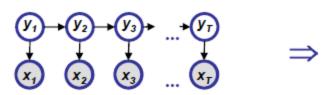
### The Sketch of Junction Tree Algorithm

#### The algorithm

- Construction of junction trees --- a special clique tree
- Propagation of probabilities --- a message-passing protocol
- Results in marginal probabilities of all cliques --- solves all queries in a single run
- A generic exact inference algorithm for any GM
- Complexity: exponential in the size of the maximal clique --a good elimination order often leads to small maximal clique,
  and hence a good (i.e., thin) JT
- Many well-known algorithms are special cases of JT
  - Forward-backward, Kalman filter, Peeling, Sum-Product ...

### A Junction Tree Algorithm for HMM

A junction tree for the HMM



Rightward pass

$$\mu_{t \to t+1}(y_{t+1}) = \sum_{y_t} \psi(y_t, y_{t+1}) \mu_{t-1 \to t}(y_t) \mu_{t\uparrow}(y_{t+1})$$

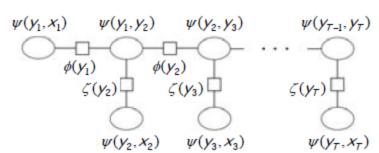
$$= \sum_{y_t} p(y_{t+1} \mid y_t) \mu_{t-1 \to t}(y_t) p(x_{t+1} \mid y_{t+1})$$

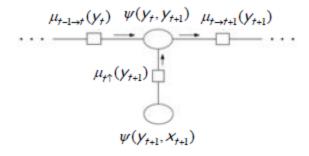
$$= p(x_{t+1} \mid y_{t+1}) \sum_{y_t} a_{y_t, y_{t+1}} \mu_{t-1 \to t}(y_t)$$

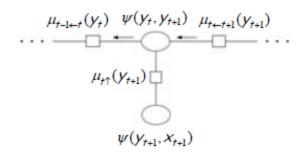
- This is exactly the forward algorithm!
- Leftward pass ...

$$\begin{split} \mu_{t-1 \leftarrow t}(y_t) &= \sum_{y_{t+1}} \psi(y_t, y_{t+1}) \mu_{t \leftarrow t+1}(y_{t+1}) \mu_{t \uparrow}(y_{t+1}) \\ &= \sum_{y_{t+1}} p(y_{t+1} \mid y_t) \mu_{t \leftarrow t+1}(y_{t+1}) p(x_{t+1} \mid y_{t+1}) \end{split}$$

This is exactly the backward algorithm!







## Summary

- Represent dependency structure with a directed acyclic graph
  - Node <-> random variable
  - Edges encode dependencies
    - Absence of edge -> conditional independence
  - Plate representation
  - A BN is a database of prob. Independence statement on variables



- The factorization theorem of the joint probability
  - Local specification → globally consistent distribution
  - Local representation for exponentially complex state-space
- Support efficient inference and learning