

# Machine Learning

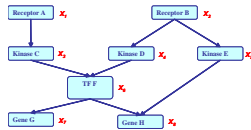
10-701/15-781, Spring 2010

## Bayesian Networks

Eric Xing

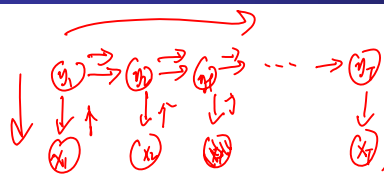
Lecture 13, March 1, 2010

Reading: Chap. 8, C.B book



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$$\begin{aligned} (1) & P(X_1 \dots X_T, Y_1 \dots Y_T) \\ &= P(X_1) P(X_2 | X_1) P(Y_1 | X_1) P(X_3 | X_2) \dots \\ & P(X_T | Y_T) \end{aligned}$$

$$\begin{aligned} (2) & P(X_t | Y_t, X_1 \dots X_{t-1}, Y_1 \dots Y_{t-1}) \\ &= P(X_t | Y_t) \\ & P(Y_t | Y_{t-1} \dots Y_1, X_t \dots X_{t-1}) \\ &= P(Y_t | Y_{t-1}) \end{aligned}$$

$$(3) P(Y_t | X_1 \dots X_T)$$

Relief Prop  
Message Passing

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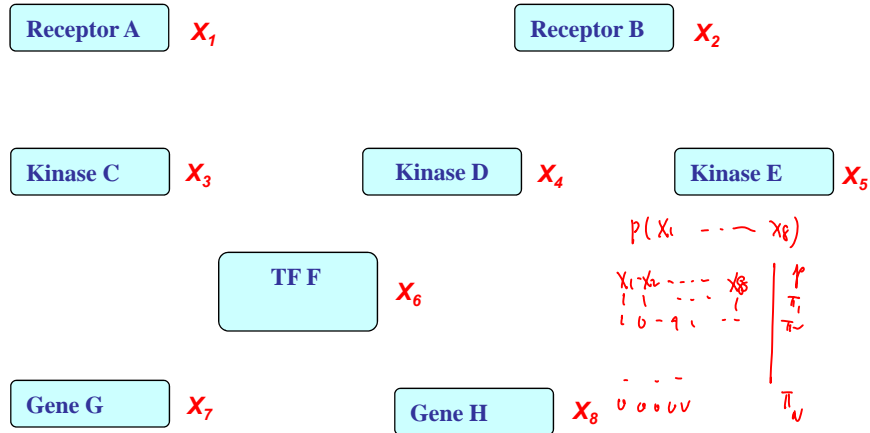
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# What is a Bayesian Network?

--- example from a signal transduction pathway



- A possible world for cellular signal transduction:



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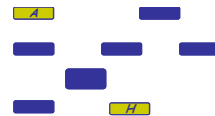
# Recap of Basic Prob. Concepts



- 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? ---  $2^8$
- 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  $p(H|A)$  would require summing over all  $2^6$  configurations of the unobserved variables

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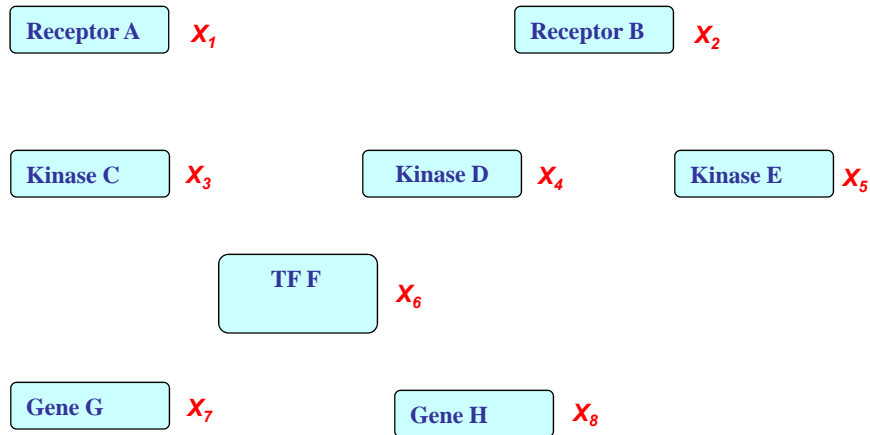
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# What is a Bayesian Network?

--- example from a signal transduction pathway



- A possible world for cellular signal transduction:



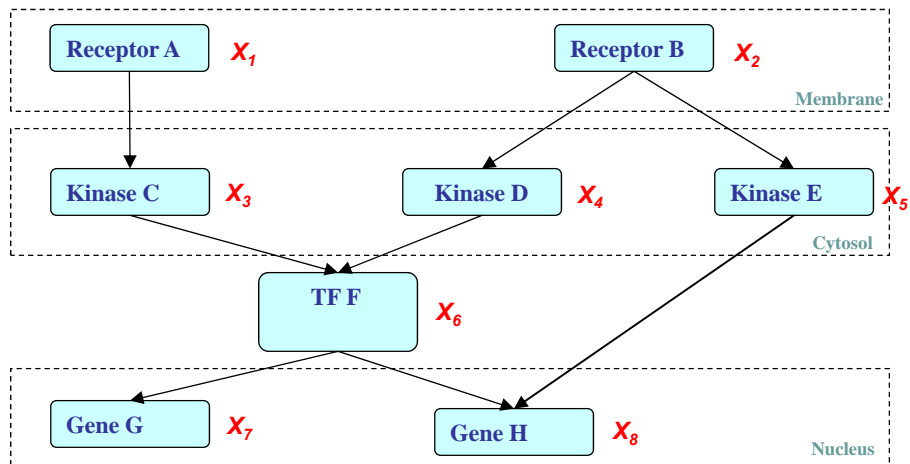
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# BN: Structure Simplifies Representation



- Dependencies among variables



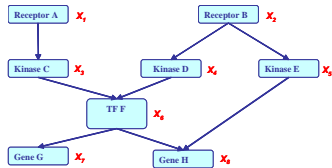
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# Bayesian Network



- 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.,



$$\begin{aligned}
 &P(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8) \\
 &= P(X_1) P(X_2) P(X_3 | X_1) P(X_4 | X_2) P(X_5 | X_2) \\
 &\quad P(X_6 | X_3, X_4) P(X_7 | X_6) P(X_8 | X_5, X_6)
 \end{aligned}$$

- Why we may favor a BN?

- Representation cost: how many probability statements are needed?
  - ~~$2 \times 2 \times 4 \times 4 \times 4 \times 8 \times 4 \times 8 = 36$~~ , an 8-fold reduction from  $2^8$ !
- Algorithms for systematic and efficient inference/learning computation
  - Exploring the graph structure and probabilistic semantics
- Incorporation of domain knowledge and causal (logical) structures

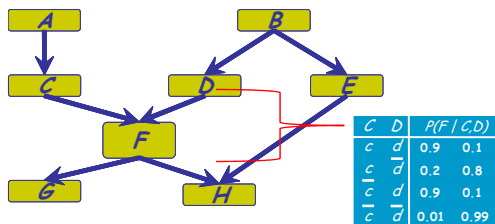
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# Specification of a BN



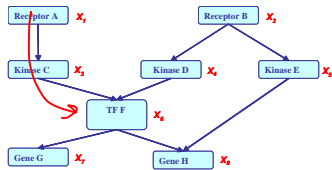
- There are two components to any GM:
  - the *qualitative* specification
  - the *quantitative* specification



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## Bayesian Network: Factorization Theorem



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

- 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”:

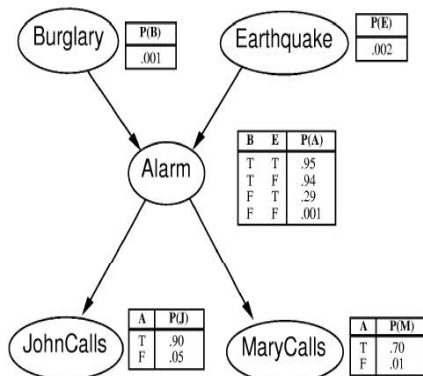
$$P(\mathbf{X}) = \prod_i P(X_i | \mathbf{X}_{\pi_i})$$

where  $\mathbf{X}_{\pi_i}$  is the set of parents of  $x_i$ .  $d$  is the number of nodes (variables) in the graph.

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## Examples



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

$$P(M|B)$$

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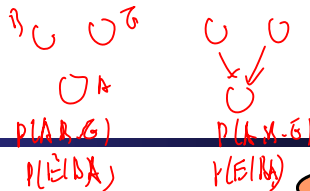
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# Qualitative Specification

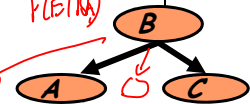


- Where does the qualitative specification come from?
  - Prior knowledge of causal relationships
  - Prior knowledge of modular relationships
  - Assessment from experts
  - Learning from data
  - We simply link a certain architecture (e.g. a layered graph)
  - ....

# Local Structures & Independencies



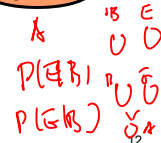
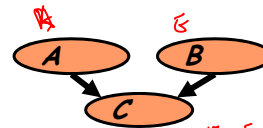
- Common parent
  - Fixing B decouples A and C  
"given the level of gene B, the levels of A and C are independent"



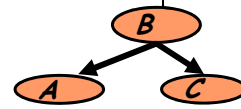
- Cascade
  - Knowing B decouples A and C  
"given the level of gene B, the level gene A provides no extra prediction value for the level of gene C"



- 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"



## A simple justification



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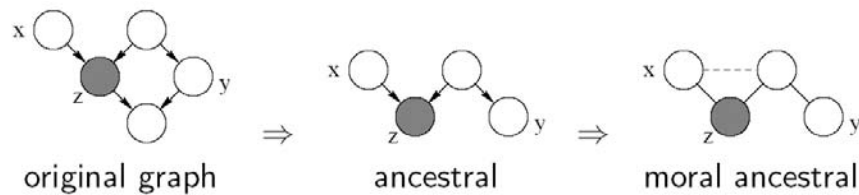
## Graph separation criterion



- 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:



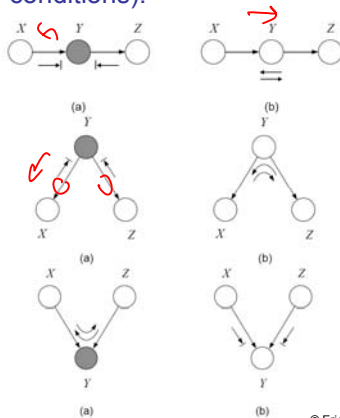
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# Global Markov properties of DAGs



- 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 below (and plus some boundary conditions):



- Defn:  $I(\theta)$  = all independence properties that correspond to d-separation:

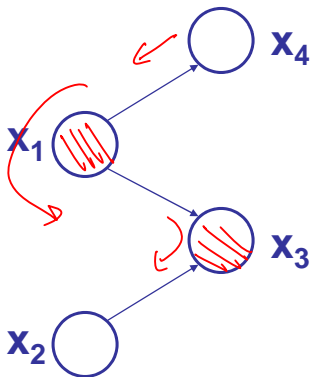
$$I(G) = \{X \perp Z | Y : \text{dsep}_G(X; Z | Y)\}$$

- D-separation is sound and complete

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## Example:



- Complete the  $I(G)$  of this graph:

$X_4 \perp X_2 | \emptyset$   
 $X_4 \perp X_3 | \emptyset$   
 $X_4 \perp X_3 | X_1$   
 $X_4 \perp X_2 | X_1, X_3$

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

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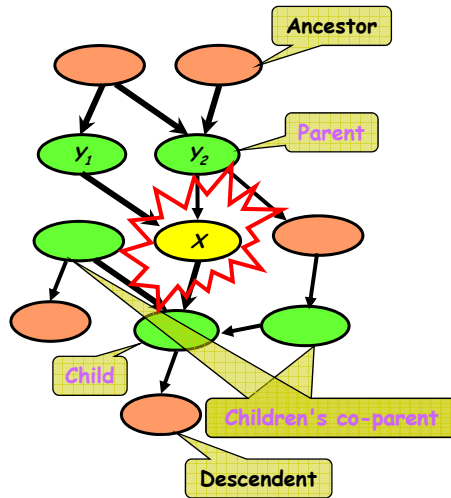


# Bayesian Network: Conditional Independence Semantics



## 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**



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# Towards quantitative specification of probability distribution

dis for  $F \rightarrow F(u, v)$   $G \rightarrow F(u, v)$



- 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  $\mathcal{D}_1$  denote the family of all distributions that satisfy  $I(G)$ ,

Let  $\mathcal{D}_2$  denote the family of all distributions that factor according to  $G$ ,

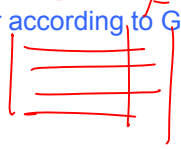
Then  $\mathcal{D}_1 \equiv \mathcal{D}_2$ .

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



$$P(A|C)P(B|C)$$

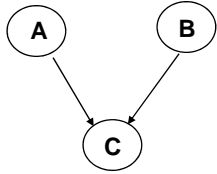
$$\neq P(A, B|C)$$



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# Quantitative Specification



$$p(A,B,C) = P(A)P(B)P(C|A,B)$$

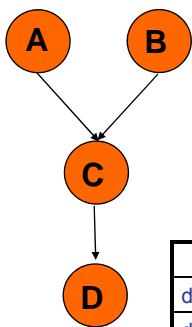
# Conditional probability tables (CPTs)



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

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

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



	a <sup>0</sup> b <sup>0</sup>	a <sup>0</sup> b <sup>1</sup>	a <sup>1</sup> b <sup>0</sup>	a <sup>1</sup> b <sup>1</sup>
c <sup>0</sup>	0.45	1	0.9	0.7
c <sup>1</sup>	0.55	0	0.1	0.3

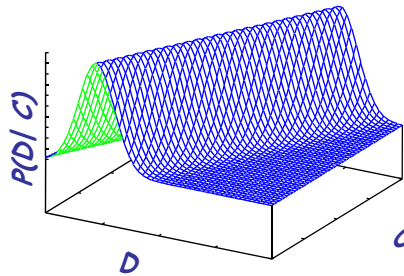
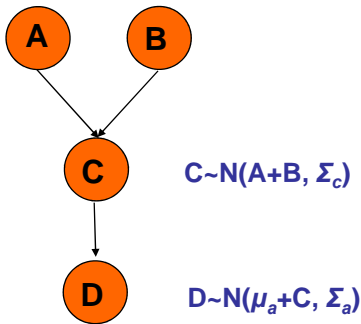
	c <sup>0</sup>	c <sup>1</sup>
d <sup>0</sup>	0.3	0.5
d <sup>1</sup>	0.7	0.5

# Conditional probability density func. (CPDs)

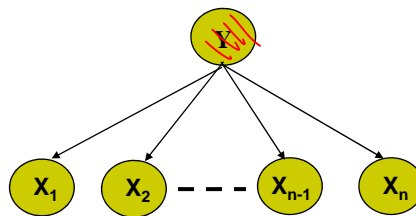


$A \sim N(\mu_a, \Sigma_a)$     $B \sim N(\mu_b, \Sigma_b)$

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



# Conditional Independencies



Label

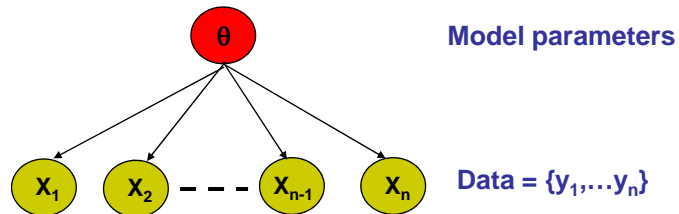
Features

What is this model

1. When Y is observed?
2. When Y is unobserved?

*NB*  
*Mixture*

# Conditionally Independent Observations



# “Plate” Notation

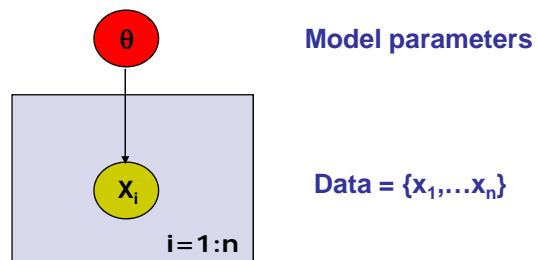
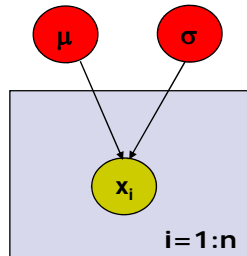


Plate = rectangle in graphical model

variables within a plate are replicated in a conditionally independent manner

## Example: Gaussian Model



Generative model:

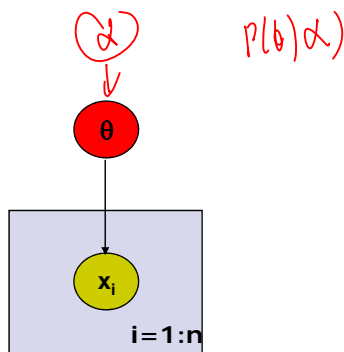
$$\begin{aligned} p(x_1, \dots, x_n \mid \mu, \sigma) &= \prod p(x_i \mid \mu, \sigma) \\ &= p(\text{data} \mid \text{parameters}) \\ &= p(D \mid \theta) \\ &\text{where } \theta = \{\mu, \sigma\} \end{aligned}$$

- Likelihood =  $p(\text{data} \mid \text{parameters})$   
=  $p(D \mid \theta)$   
=  $L(\theta)$
- Likelihood tells us how likely the observed data are conditioned on a particular setting of the parameters
  - Often easier to work with  $\log L(\theta)$

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## Bayesian models



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## Example: modeling text



### A Hierarchical Phrase-Based Model for Statistical Machine Translation

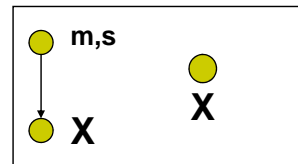
We present a statistical phrase-based Translation model that uses *hierarchical phrases*—phrases that contain sub-phrases. The model is formally a synchronous context-free grammar but is learned from a bitext without any syntactic information. Thus it can be seen as a shift to the *formal machinery* of syntax based translation systems without any *linguistic* commitment. In our experiments using BLEU as a metric, the hierarchical Phrase based model achieves a relative improvement of 7.5% over Pharaoh, a state-of-the-art phrase-based system.

## More examples



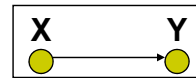
### Density estimation

Parametric and nonparametric methods



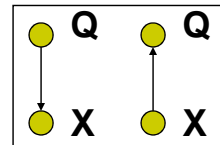
### Regression

Linear, conditional mixture, nonparametric



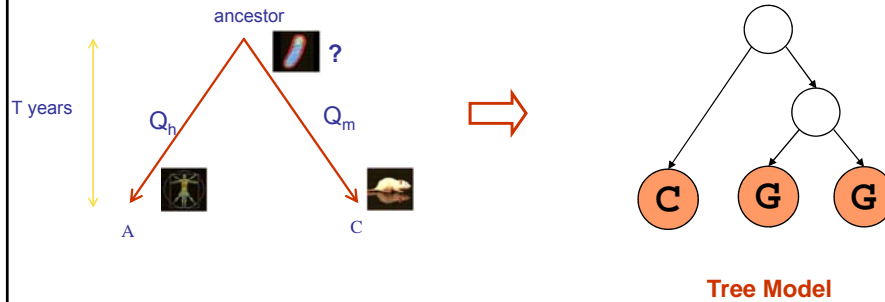
### Classification

Generative and discriminative approach



## Example, con'd

- Evolution

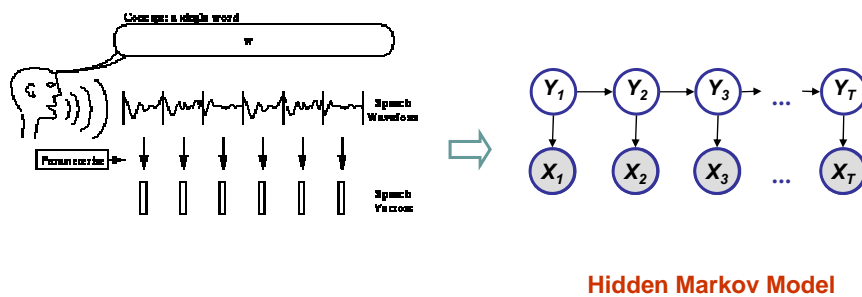


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## Example, con'd

- Speech recognition

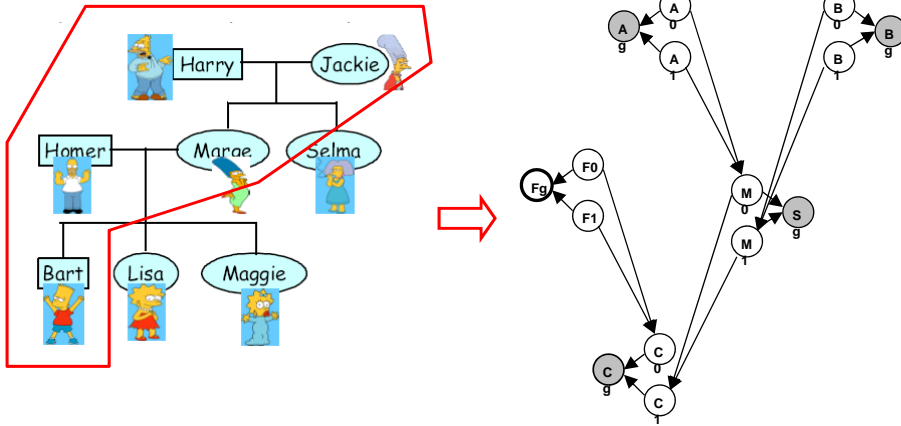


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# Example, con'd

- Genetic Pedigree

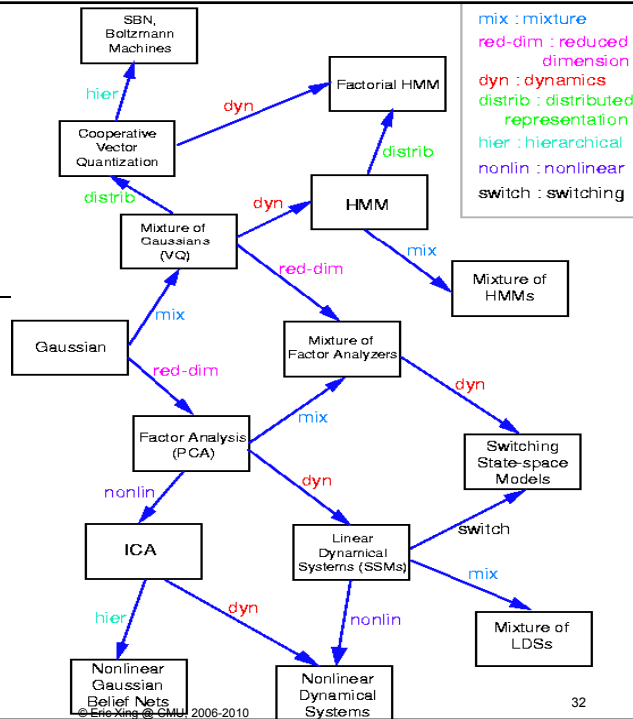


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# An (incomplete) genealogy of BNs

(Picture by Zoubin Ghahramani and Sam Roweis)



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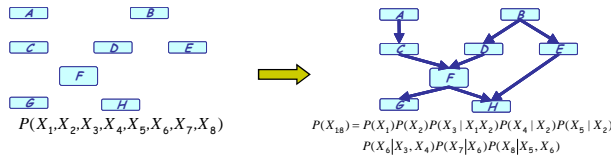
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# BN and Graphical Models



- 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



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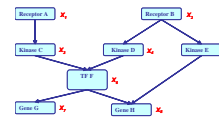
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# Two types of GMs



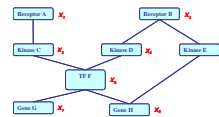
- **Directed edges** give **causality** relationships (Bayesian Network or Directed Graphical Model):

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



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

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



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# 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  $X=\text{true}$  if ( $Y=\text{false}$  and  $Z=\text{true}$ )?
  - What is the joint distribution of  $(X, Y)$  if  $Z=\text{false}$ ?
  - 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

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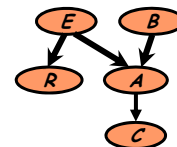
# Learning BNs (or GMs)



## The goal:

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

$(B, E, A, C, R) = (T, F, F, T, F)$   
 $(B, E, A, C, R) = (T, F, T, T, F)$   
 .....  
 $(B, E, A, C, R) = (F, T, T, T, F)$



$E$	$B$	$P(A   E, B)$	
$e$	$b$	0.9	0.1
$e$	$\bar{b}$	0.2	0.8
$\bar{e}$	$b$	0.9	0.1
$\bar{e}$	$\bar{b}$	0.01	0.99

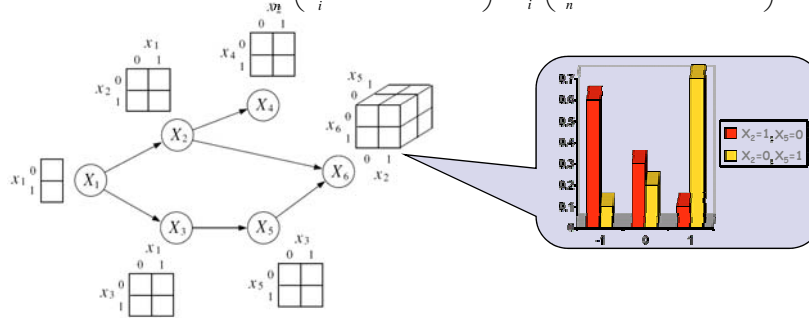
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## MLE for general BN parameters

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

$$\mathcal{L}(\theta; D) = \log p(D | \theta) = \log \prod_i \left( \prod_{i'} p(x_{n,i'} | \mathbf{x}_{n,\pi_i}, \theta_i) \right) = \sum_i \left( \sum_{n'} \log p(x_{n,i'} | \mathbf{x}_{n,\pi_i}, \theta_i) \right)$$



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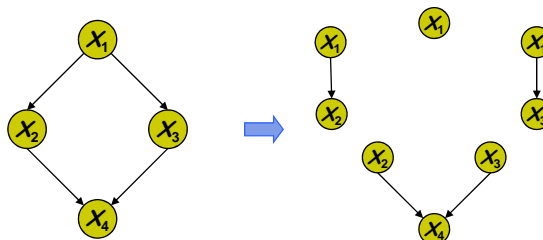
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## Example: decomposable likelihood of a directed model

- Consider the distribution defined by the directed acyclic GM:

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

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



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## E.g.: MLE 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,  $\mathbf{x}_{\pi_i}$  will have a composite state, and the CPD will be a high-dimensional table
- The sufficient statistics are counts of family configurations

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

- The log-likelihood is  $\ell(\theta; \mathcal{D}) = \log \prod_{i,j,k} \theta_{ijk}^{n_{ijk}} = \sum_{i,j,k} n_{ijk} \log \theta_{ijk}$

- Using a Lagrange multiplier to enforce  $\sum_j \theta_{ijk} = 1$ , we get:

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

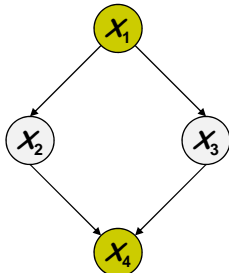


## What if some nodes are not observed?



- Consider the distribution defined by the directed acyclic GM:

$$p(x \mid \theta) = p(x_1 \mid \theta_1) p(x_2 \mid x_1, \theta_2) p(x_3 \mid x_1, \theta_3) p(x_4 \mid x_2, x_3, \theta_4)$$



- Need to compute  $p(x_H \mid x_V) \rightarrow$  inference

# Summary



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

