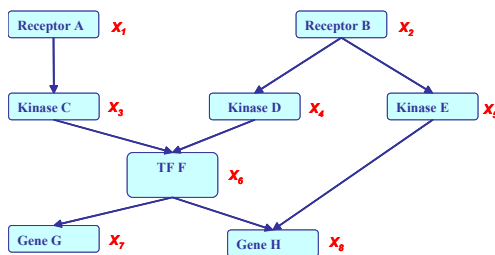
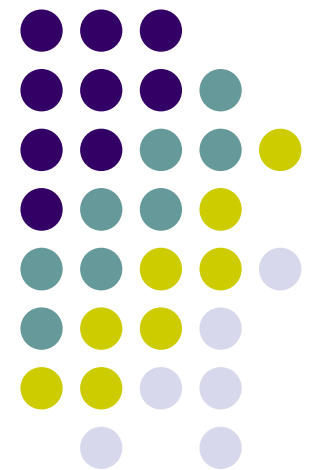


Machine Learning

10-701, Fall 2016

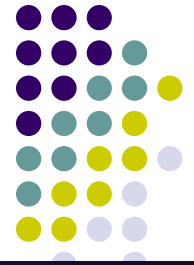
Graphical Models and Exact Inference

Eric Xing



Lecture 17, November 7, 2016

Reading: Chap. 8, C.B book

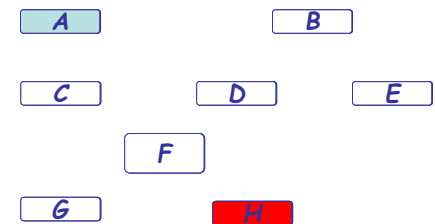


Recap of Basic Prob. Concepts

- 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?**



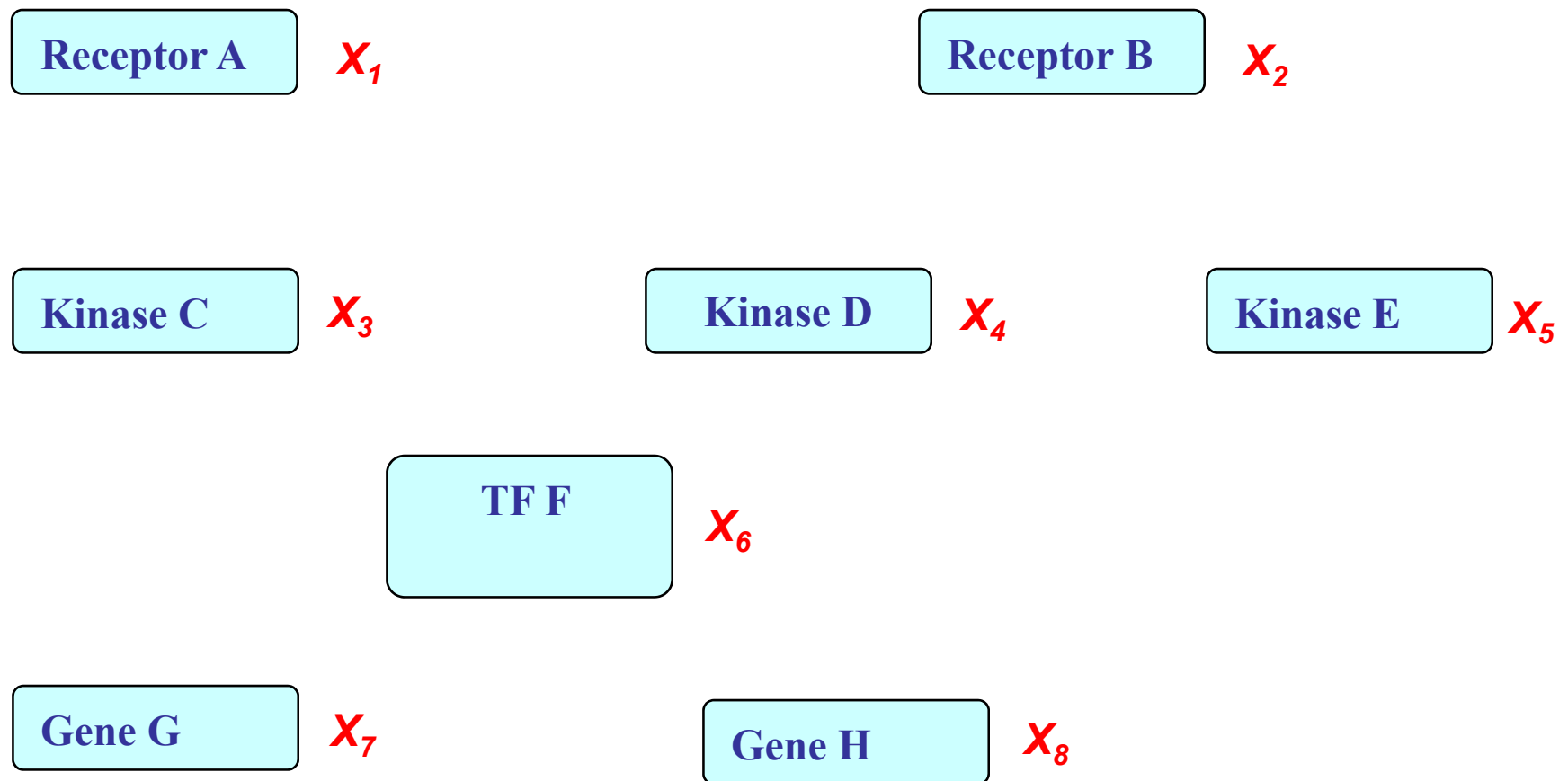
- Learning: where do we get all this probabilities?
 - Maximal-likelihood estimation? but how many data do we need?
 - Are there other est. principles?
 - Where do we put domain knowledge in terms of plausible relationships between variables, and plausible values of the probabilities?
- Inference: If 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

What is a Graphical Model?

--- Multivariate Distribution in High-D Space



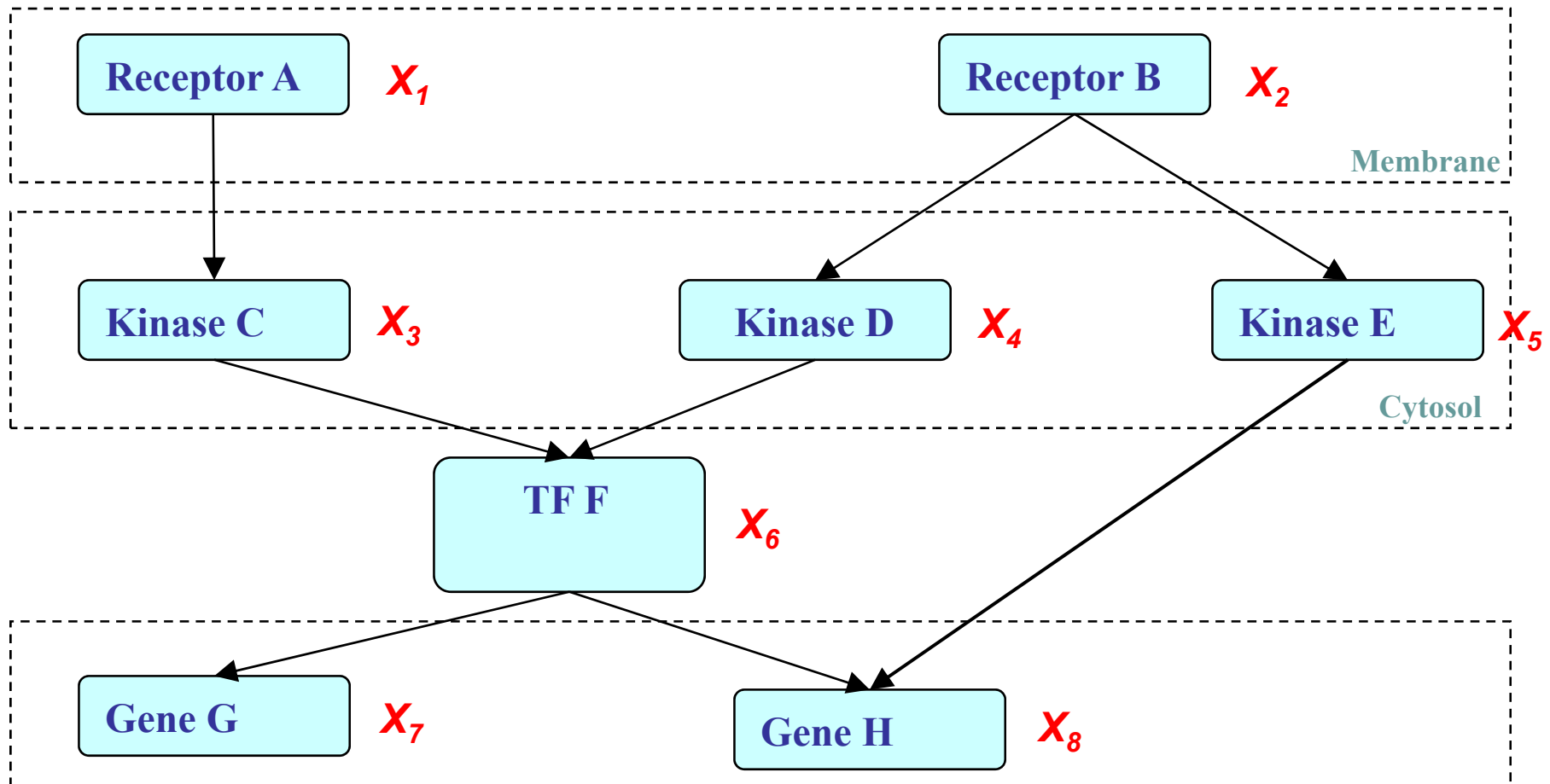
- A possible world for cellular signal transduction:



GM: Structure Simplifies Representation



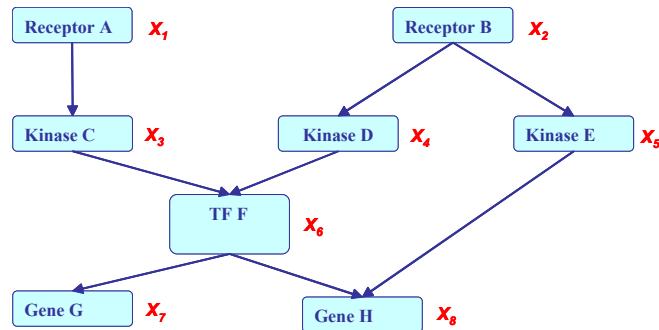
- Dependencies among variables





Probabilistic Graphical Models

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



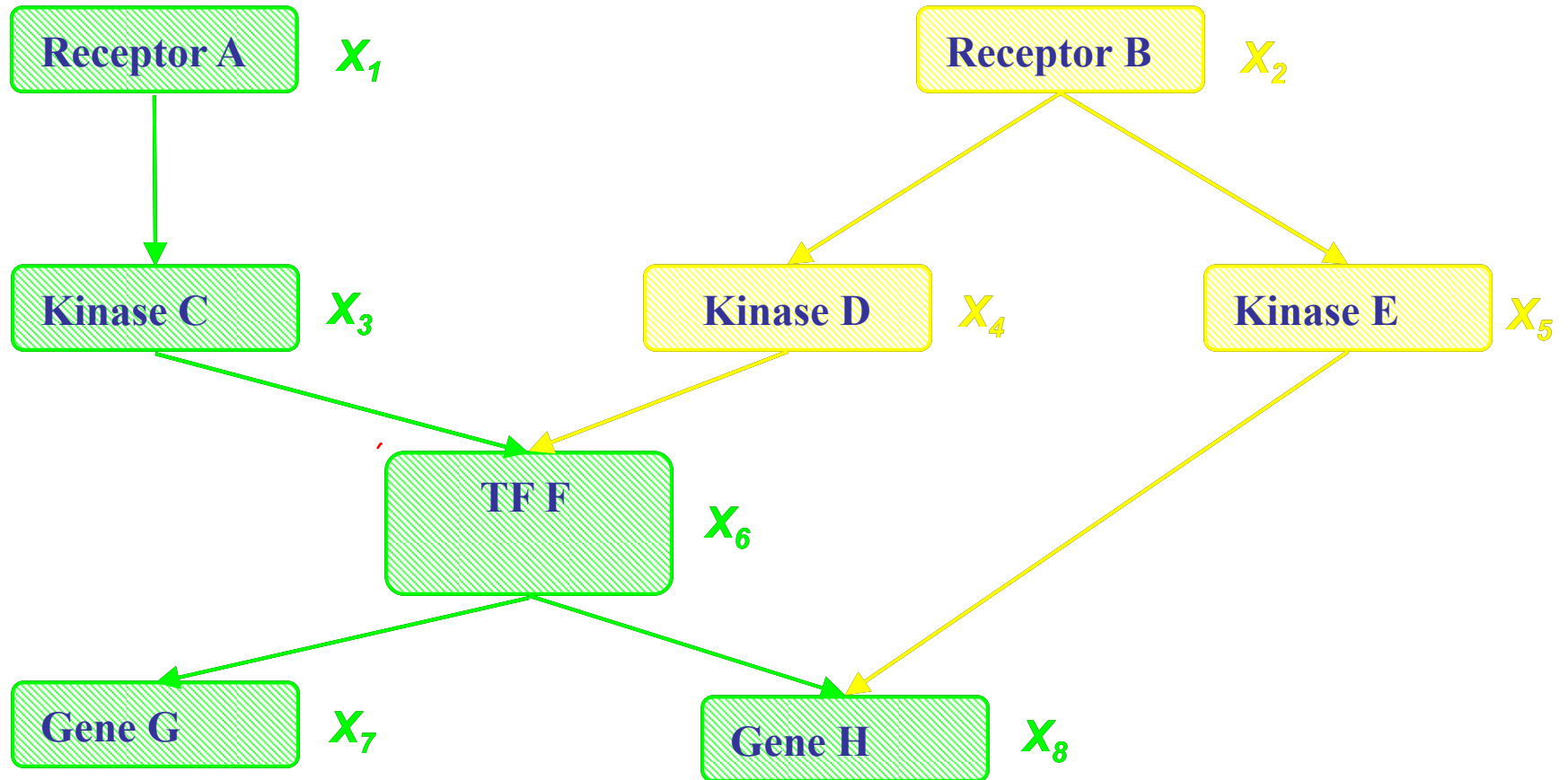
$$\begin{aligned} &P(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8) \\ &= P(X_1) P(X_2) P(X_3/X_1) P(X_4/X_2) P(X_5/X_2) \\ &P(X_6/X_3, X_4) P(X_7/X_6) P(X_8/X_5, X_6) \end{aligned}$$

Stay tune for what are these independencies!

- Why we may favor a PGM?
 - Incorporation of domain knowledge and causal (logical) structures
1+1+2+2+2+4+2+4=18, a 16-fold reduction from 2^8 in representation cost !



GM: Data Integration



More Data Integration



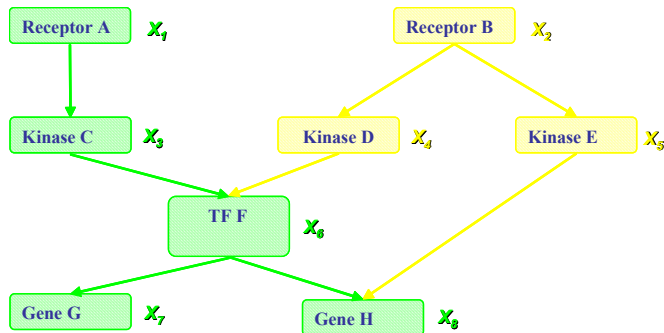
- Text + Image + Network → Holistic Social Media

- Genome + Proteome + Transcriptome + Phenome + ... → PanOmic Biology



Probabilistic Graphical Models

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- Why we may favor a PGM?
 - Incorporation of domain knowledge and causal (logical) structures
 $2+2+4+4+4+8+4+8=36$, an 8-fold reduction from 2^8 in representation cost !
 - Modular combination of heterogeneous parts – data fusion

Rational Statistical Inference



The Bayes Theorem:

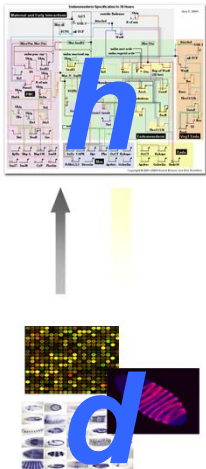
Posterior probability

Likelihood

Prior probability

$$p(h | d) = \frac{p(d | h) p(h)}{\sum_{h' \in H} p(d | h') p(h')}$$

Sum over space of hypotheses

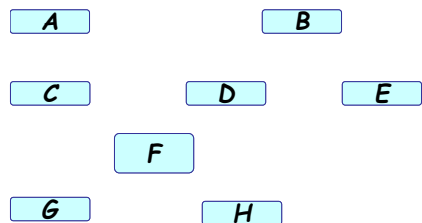


- This allows us to capture uncertainty about the model in a principled way
- But how can we specify and represent a complicated model?
 - Typically the number of genes need to be modeled are in the order of thousands!



GM: MLE and Bayesian Learning

- Probabilistic statements of Θ is conditioned on the values of the observed variables \mathbf{A}_{obs} and prior $p(\cdot | \chi)$

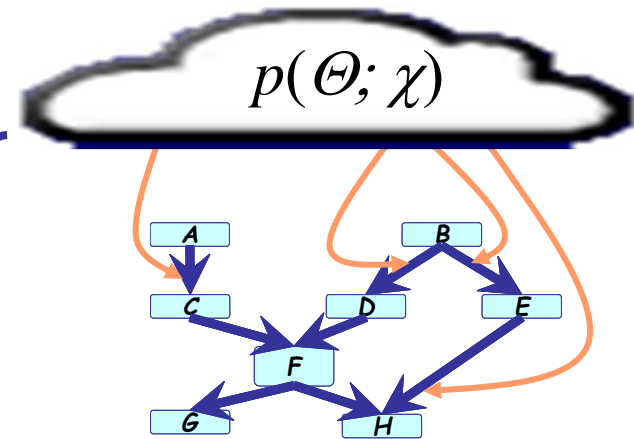
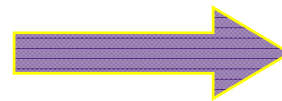


$(A,B,C,D,E,\dots)=(T,F,F,T,F,\dots)$

$\mathbf{A} = (A,B,C,D,E,\dots)=(T,F,T,T,F,\dots)$

.....

$(A,B,C,D,E,\dots)=(F,T,T,T,F,\dots)$



C	D	$P(F C,D)$	
c	d	0.9	0.1
c	\bar{d}	0.2	0.8
\bar{c}	d	0.9	0.1
\bar{c}	\bar{d}	0.01	0.99

$$\Theta_{\text{Bayes}} = \int \Theta p(\Theta | \mathbf{A}, \chi) d\Theta$$

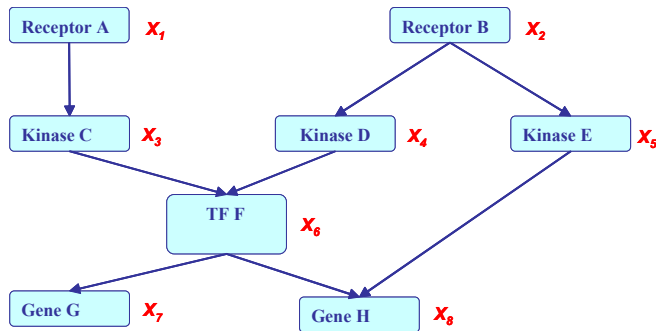
$$p(\Theta | \mathbf{A}; \chi) \propto p(\mathbf{A} | \Theta) p(\Theta; \chi)$$

posterior
likelihood
prior



Probabilistic Graphical Models

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



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- Why we may favor a PGM?
 - Incorporation of domain knowledge and causal (logical) structures
 $2+2+4+4+4+8+4+8=36$, an 8-fold reduction from 2^8 in representation cost !
 - Modular combination of heterogeneous parts – data fusion
 - Bayesian Philosophy

- Knowledge meets data





So What Is a PGM After All?

In a nutshell:

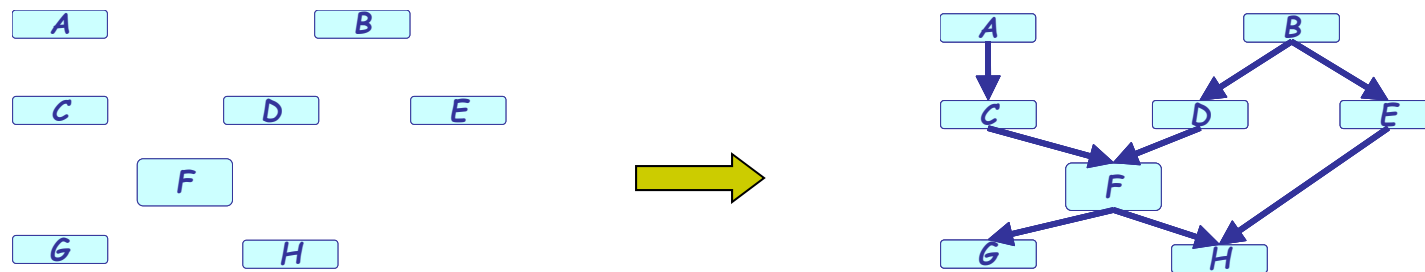
PGM = Multivariate Statistics + Structure

GM = Multivariate Obj. Func. + Structure



So What Is a PGM After All?

- The informal blurb:
 - 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**



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

$$P(X_{1:8}) = P(X_1)P(X_2)P(X_3 | X_1 X_2)P(X_4 | X_2)P(X_5 | X_2) \\ P(X_6 | X_3, X_4)P(X_7 | X_6)P(X_8 | X_5, X_6)$$

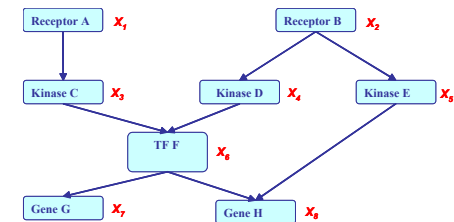
- A more formal description:
 - It refers to a family of distributions on a set of random variables that are compatible with all the probabilistic independence propositions encoded by a graph that connects these variables



Two types of GMs

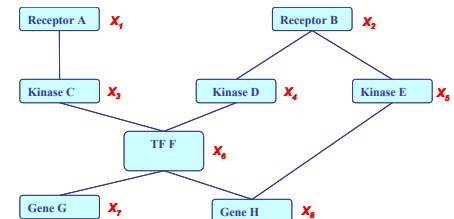
- Directed edges give causality relationships (Bayesian Network or Directed Graphical Model):

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



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

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



Towards structural 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**

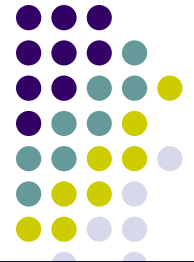
For a graph G ,

Let \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 ,

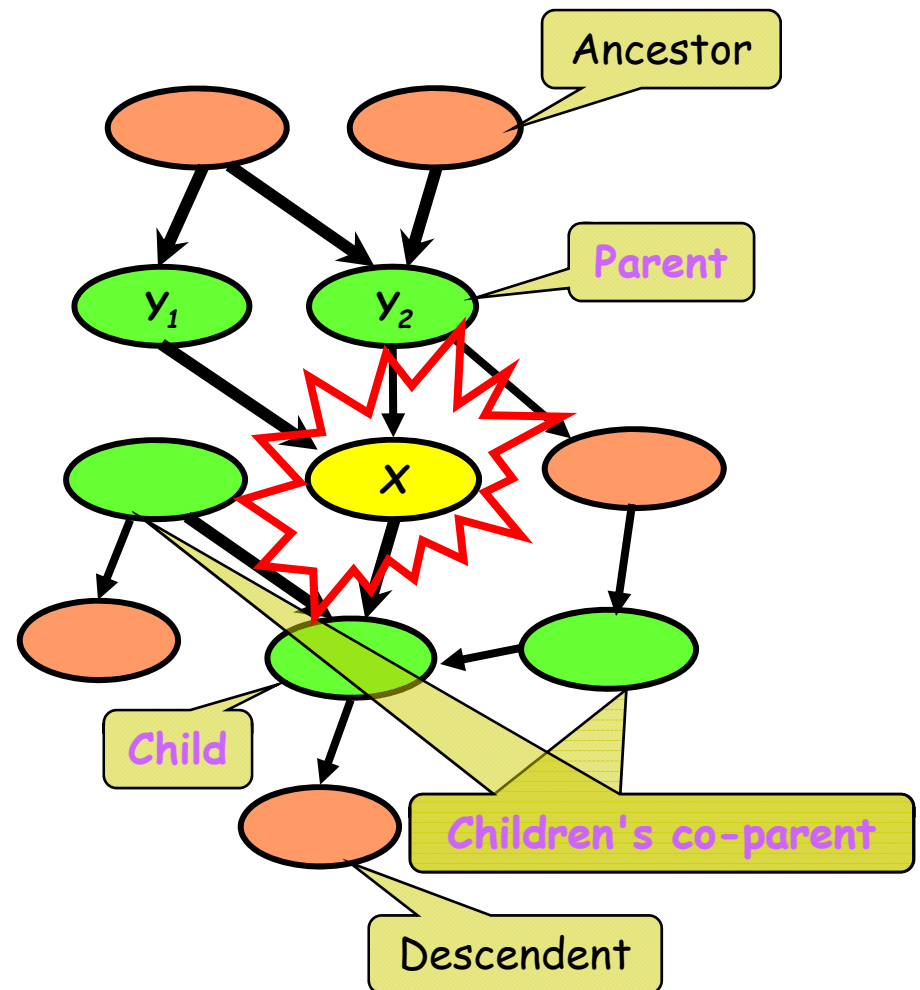
Then $\mathcal{D}_1 \equiv \mathcal{D}_2$.

Bayesian Networks



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

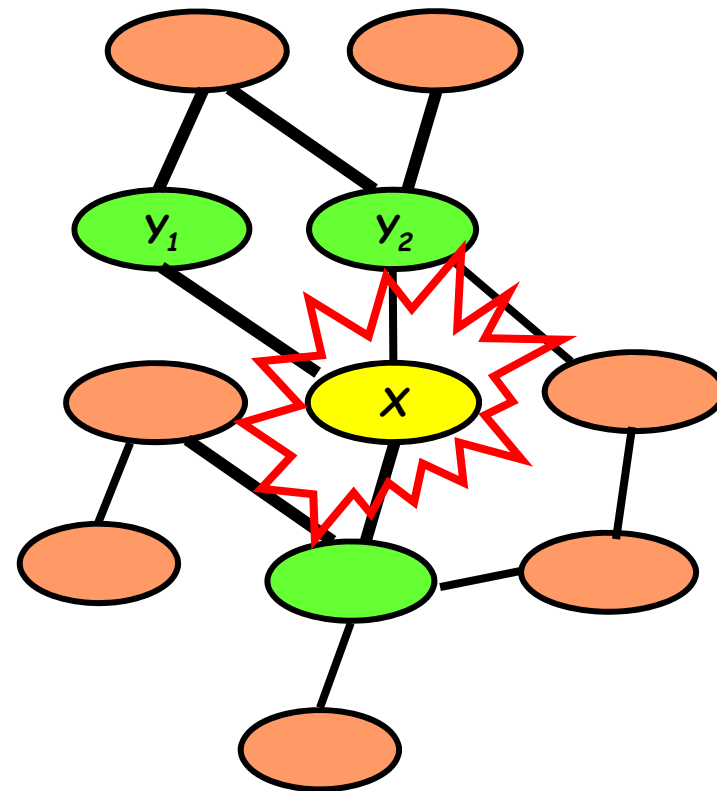




Markov Random Fields

Structure: *undirected graph*

- Meaning: a node is **conditionally independent** of every other node in the network given its **Directed neighbors**
- Local contingency functions (**potentials**) and the **cliques** in the graph completely determine the **joint** dist.
- Give **correlations** between variables, but no explicit way to generate samples

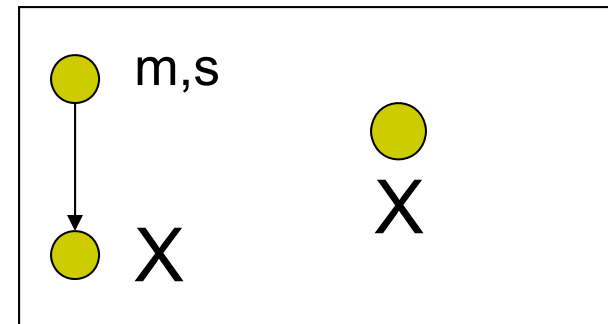




GMs are your old friends

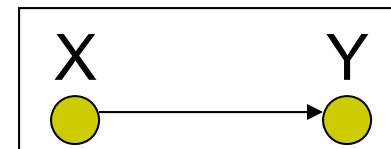
Density estimation

Parametric and nonparametric methods



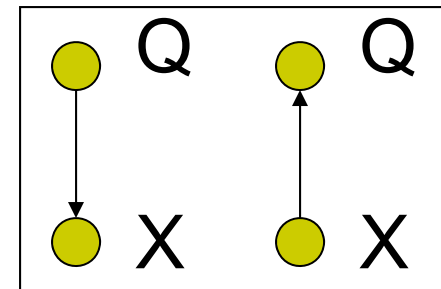
Regression

Linear, conditional mixture, nonparametric



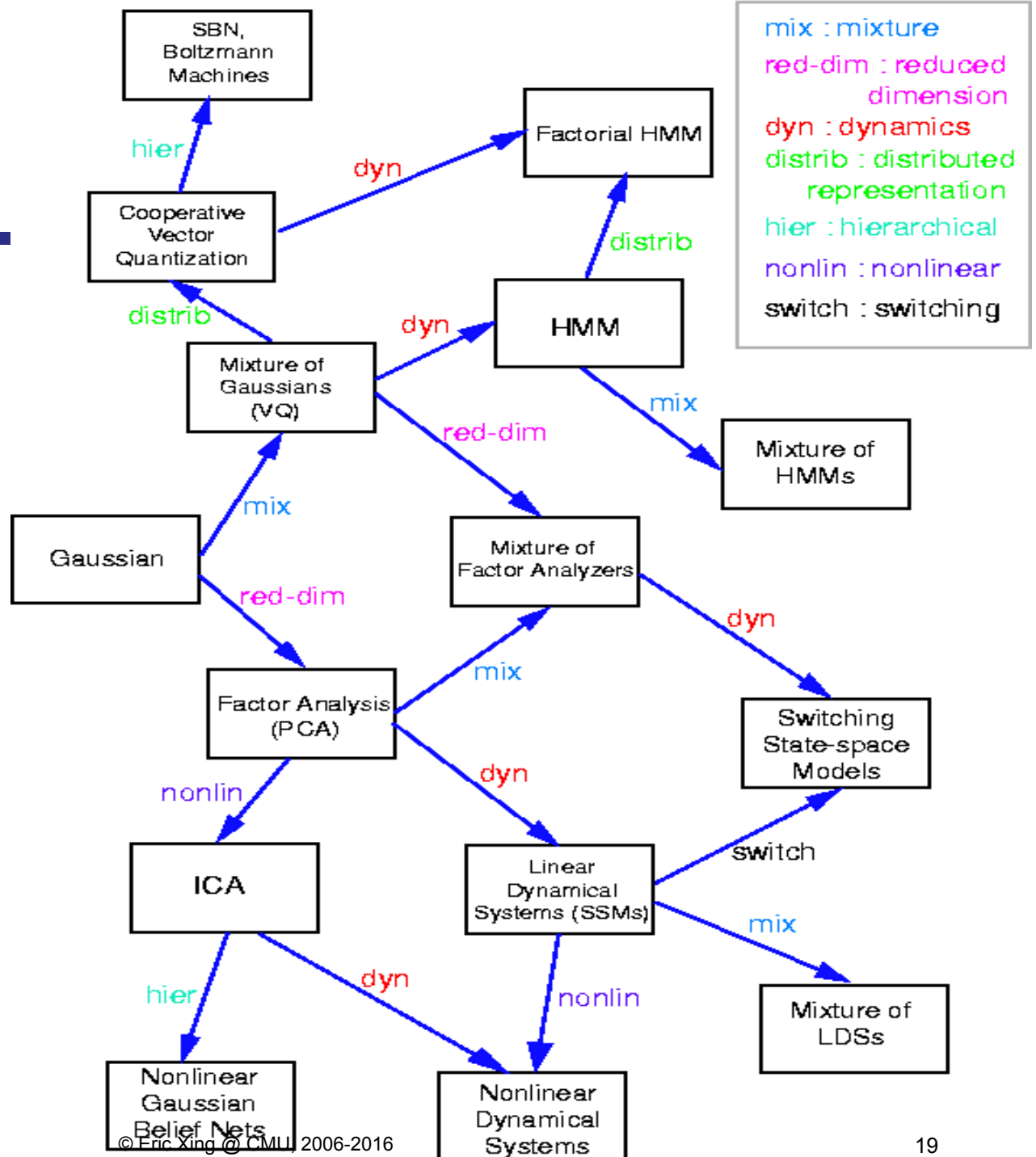
Classification

Generative and discriminative approach



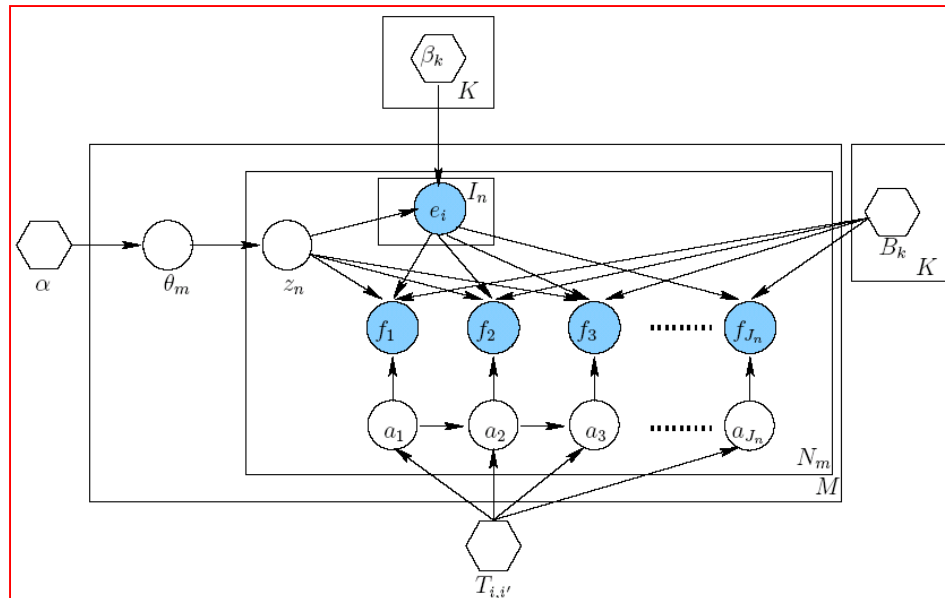
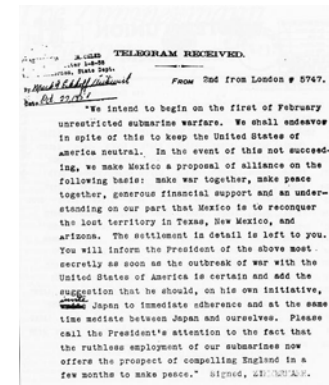
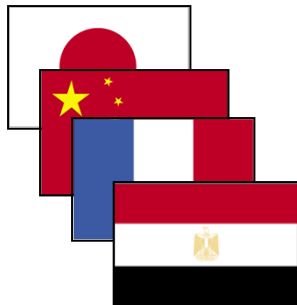
Clustering

An (incomplete) genealogy of graphical models



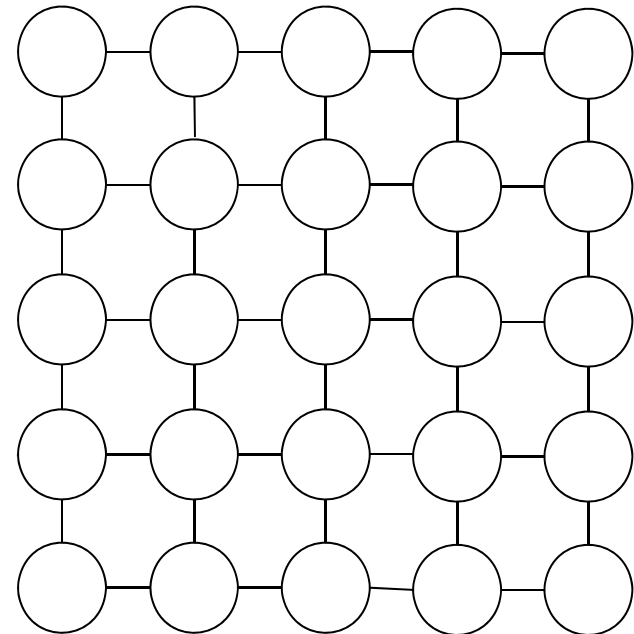
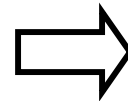
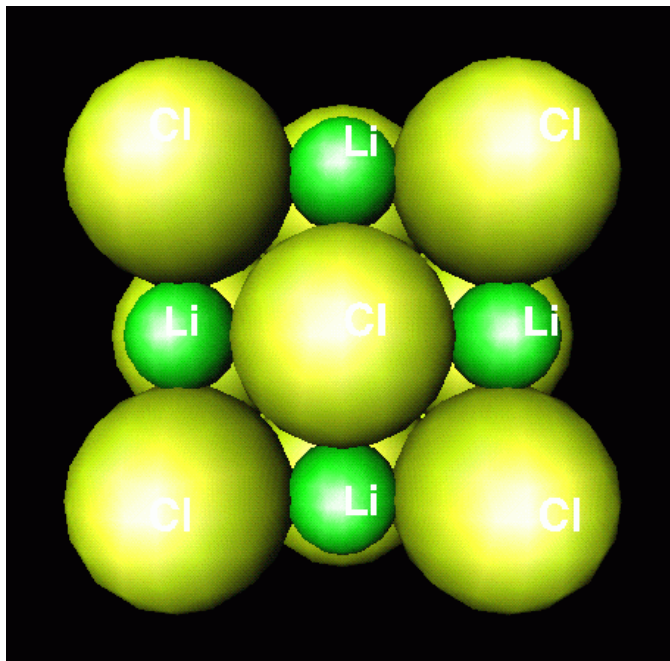
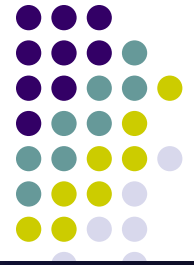
(Picture by Zoubin Ghahramani and Sam Roweis)

Fancier GMs: machine translation



The HM-BiTAM model
(B. Zhao and E.P Xing,
ACL 2006)

Fancier GMs: solid state physics



Ising/Potts model

A Generative Scheme for model design



Why graphical models



- A language for communication
 - A language for computation
 - A language for development
-
- Origins:
 - Wright 1920's
 - Independently developed by Spiegelhalter and Lauritzen in statistics and Pearl in computer science in the late 1980's

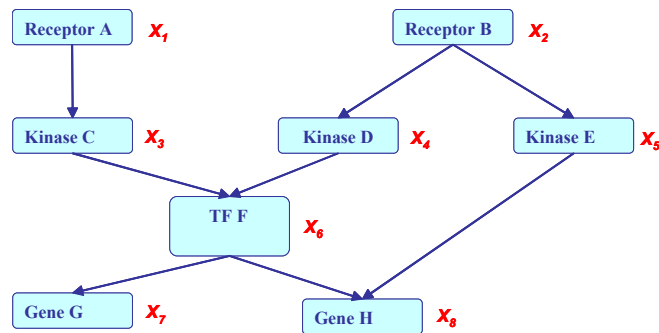


Why graphical models

- **Probability theory** provides the **glue** whereby the parts are combined, ensuring that the system as a whole is consistent, and providing ways to interface models to data.
- The **graph theoretic** side of graphical models provides both an intuitively appealing interface by which humans can model highly-interacting sets of variables as well as a data structure that lends itself naturally to the design of efficient general-purpose algorithms.
- **Many of the classical multivariate probabilistic systems** studied in fields such as statistics, systems engineering, information theory, pattern recognition and statistical mechanics **are special cases of the general graphical model formalism**
- The graphical model framework provides a way to view all of these systems as instances of a **common underlying formalism**.



Bayesian Network: Factorization Theorem



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

- **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”:

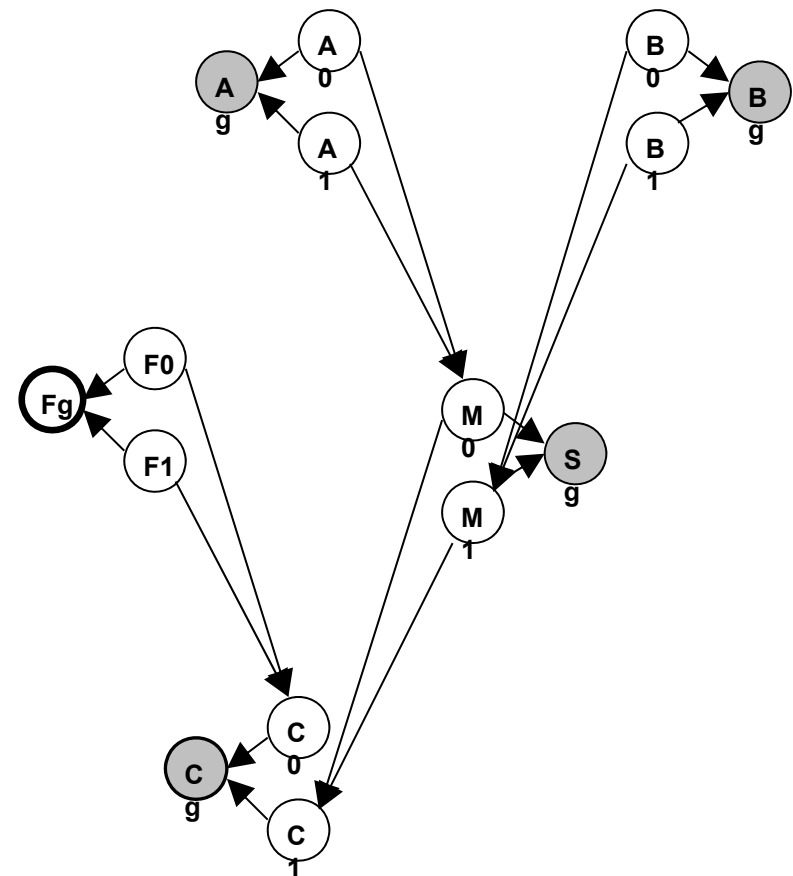
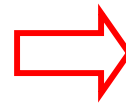
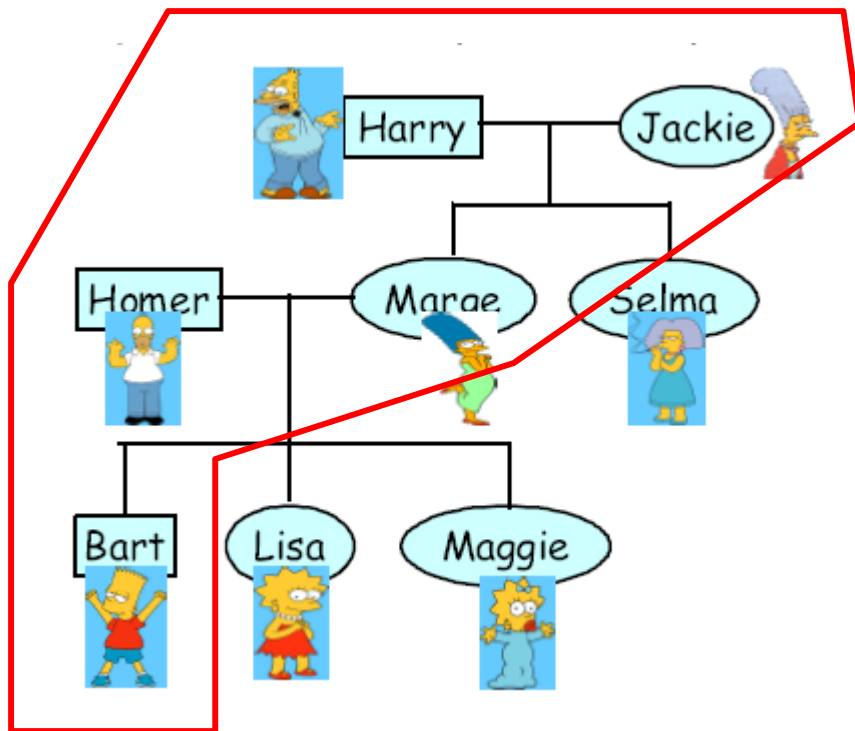
$$P(\mathbf{X}) = \prod_i P(X_i | \mathbf{X}_{\pi_i})$$

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



Example: a pedigree of people

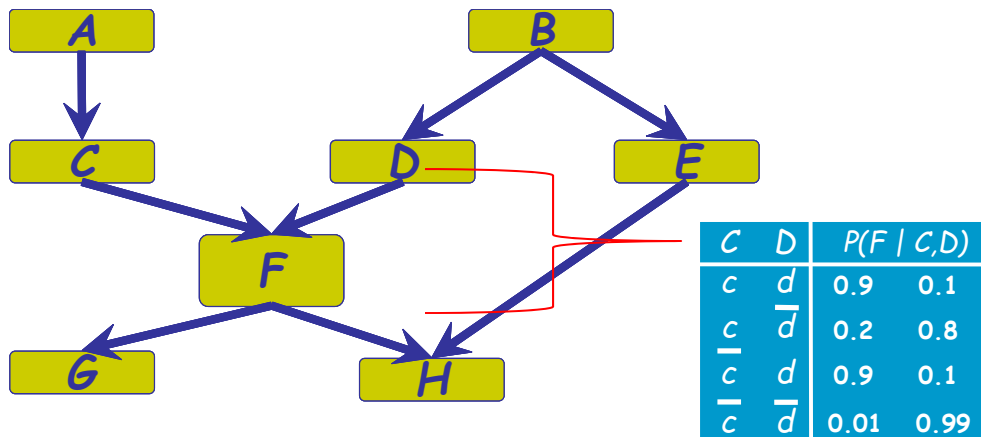
- Genetic Pedigree





Specification of a BN

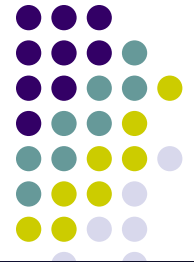
- There are two components to any GM:
 - the *qualitative* specification
 - the *quantitative* specification



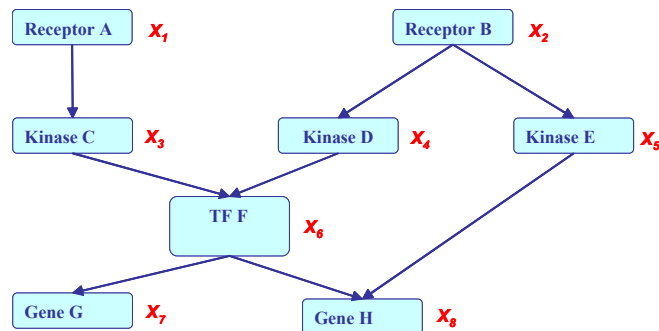
Qualitative Specification



- 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)
 - ...



Bayesian Network: Factorization Theorem



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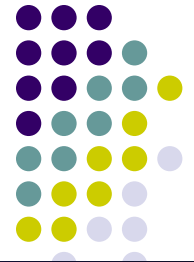
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where \mathbf{X}_{π_i} is the set of parents of x_i . d is the number of nodes (variables) in the graph.

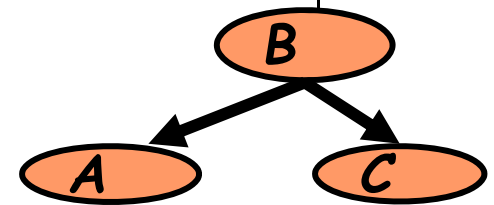
Local Structures & Independencies



- 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

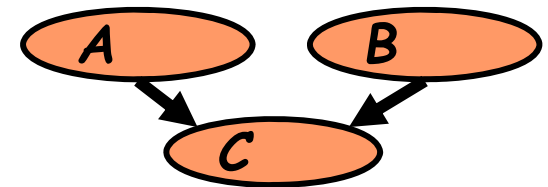
"given the level of gene B, the level gene A provides no extra prediction value for the level of gene C"



- V-structure

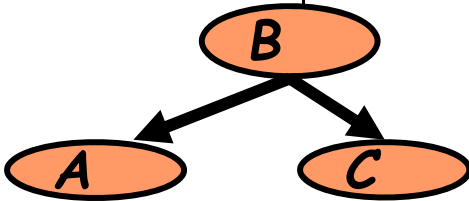
- Knowing C couples A and B because A can "explain away" B w.r.t. C

"If A correlates to C, then chance for B to also correlate to B will decrease"



- The language is compact, the concepts are rich!

A simple justification



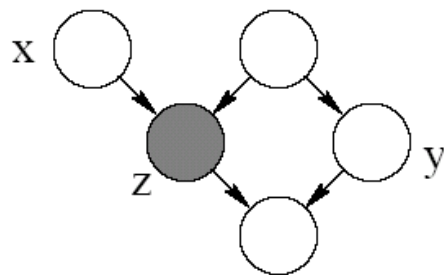


Graph separation criterion

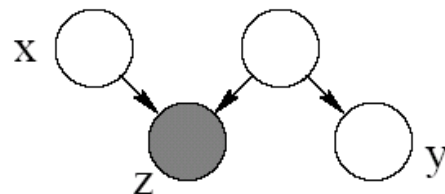
- D-separation criterion for Bayesian networks (D for Directed edges):

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

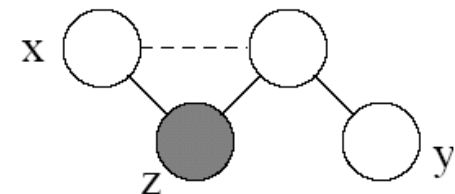
- Example:



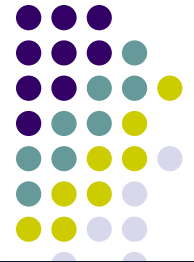
original graph



ancestral



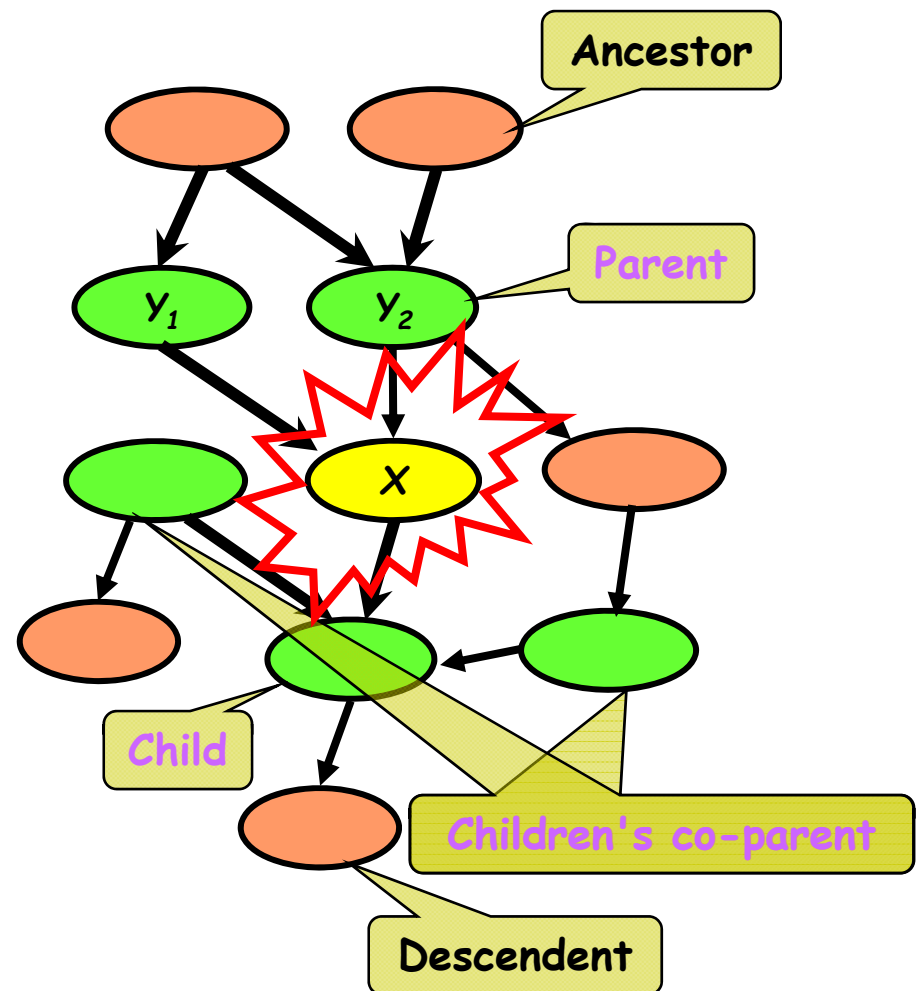
moral ancestral



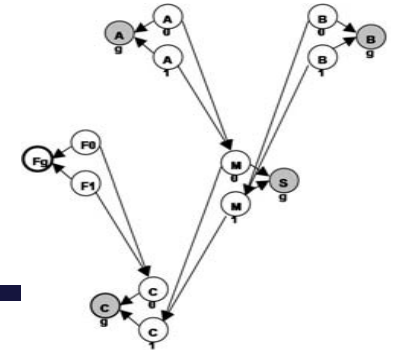
Local Markov properties of DAGs

Structure: DAG

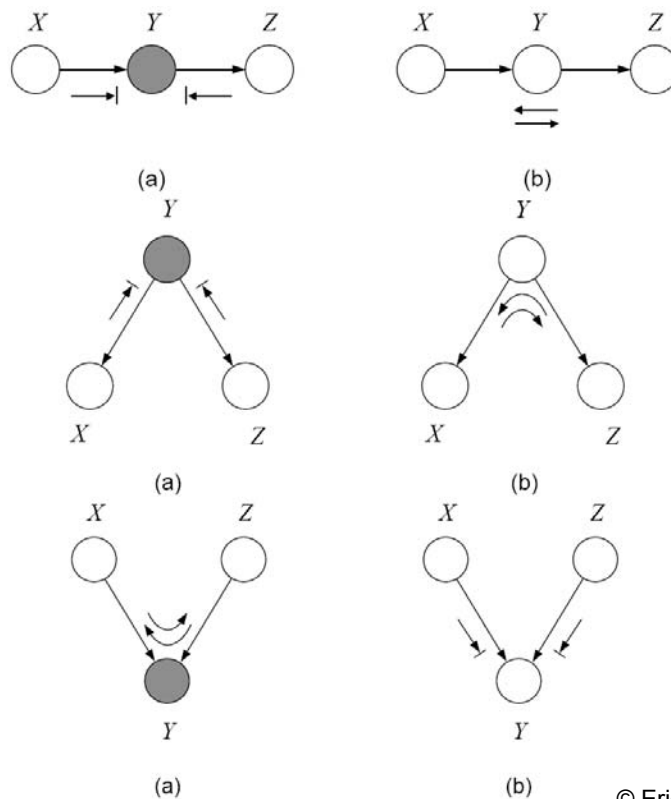
- Meaning: a node is **conditionally independent** of every other node in the network outside its **Markov blanket**
- Local conditional distributions (**CPD**) and the **DAG** completely determine the **joint** dist.
- Give **causality** relationships, and facilitate a **generative** process



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):



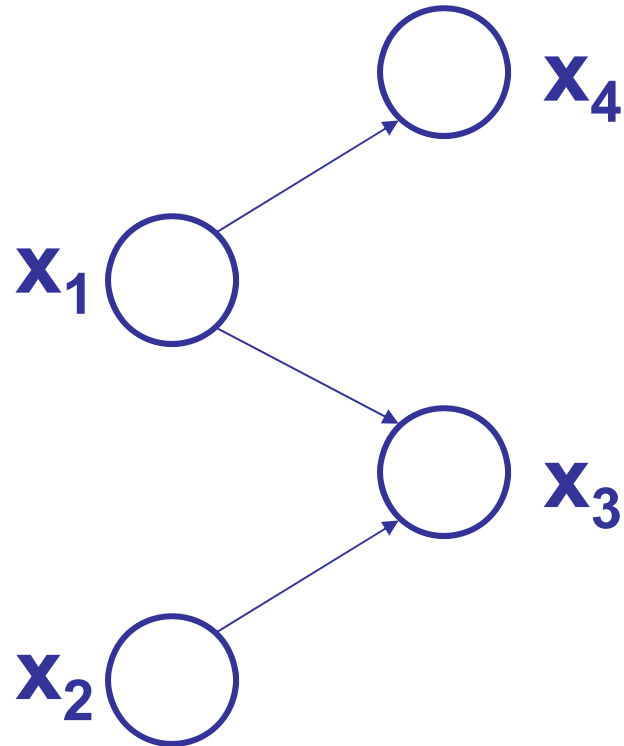
- **Defn:** $I(G)$ = all independence properties that correspond to d-separation:

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

- **D-separation is sound and complete**



Example:



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

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

Towards 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**

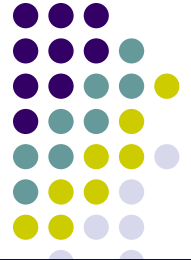
For a graph G ,

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

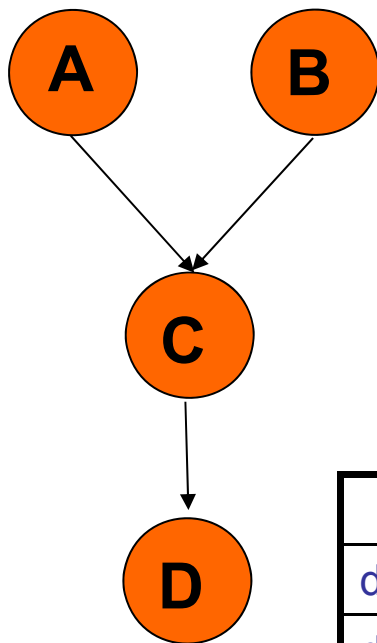
Conditional probability tables (CPTs)



a^0	0.75
a^1	0.25

b^0	0.33
b^1	0.67

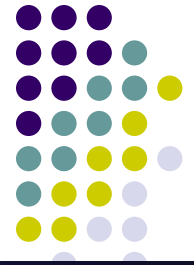
$$P(a,b,c,d) = P(a)P(b)P(c|a,b)P(d|c)$$



	a^0b^0	a^0b^1	a^1b^0	a^1b^1
c^0	0.45	1	0.9	0.7
c^1	0.55	0	0.1	0.3

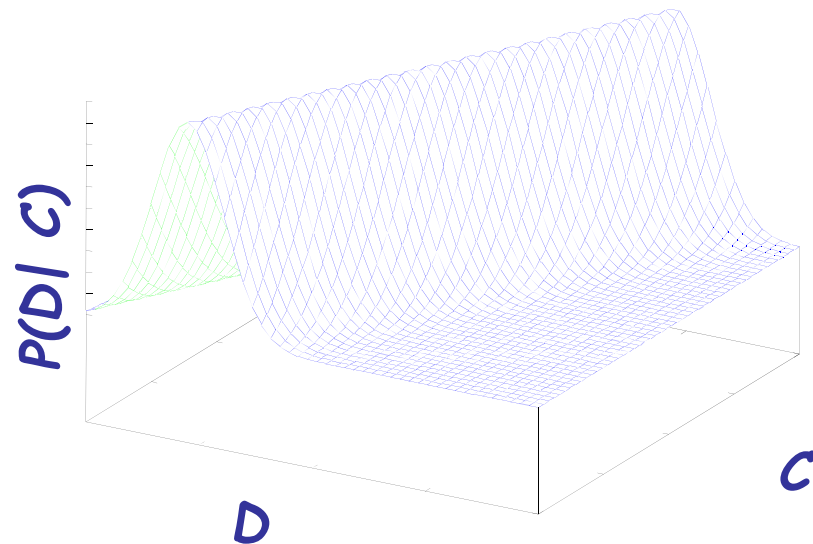
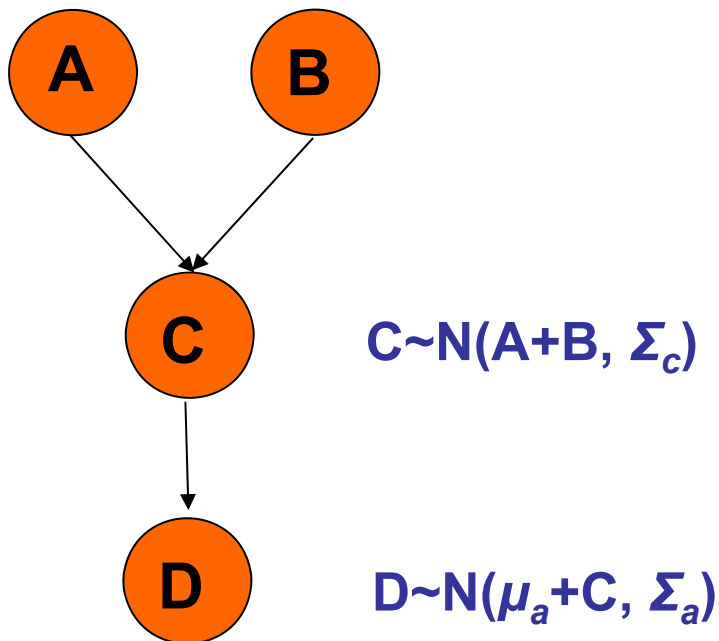
	c^0	c^1
d^0	0.3	0.5
d^1	0.7	0.5

Conditional probability density func. (CPDs)

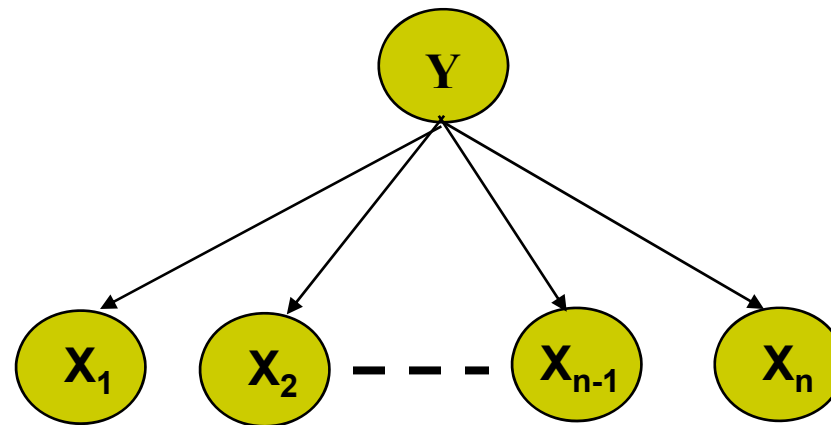
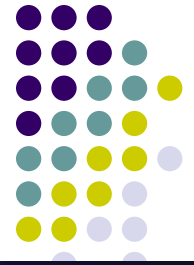


$$A \sim N(\mu_a, \Sigma_a) \quad B \sim N(\mu_b, \Sigma_b)$$

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



Conditional Independencies



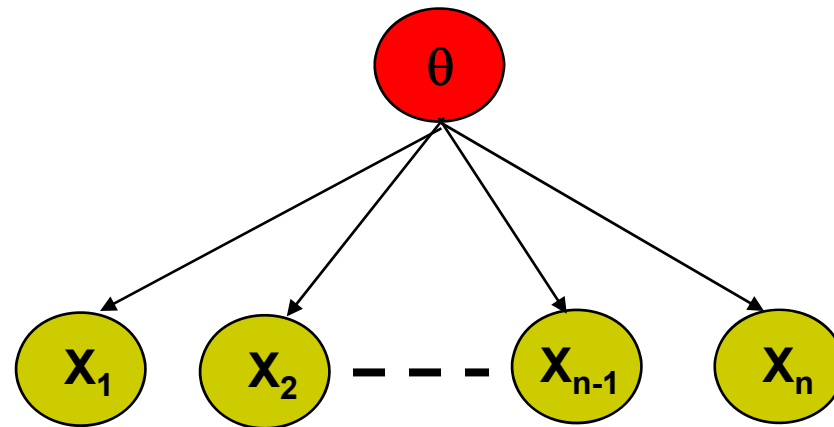
Label

Features

What is this model

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

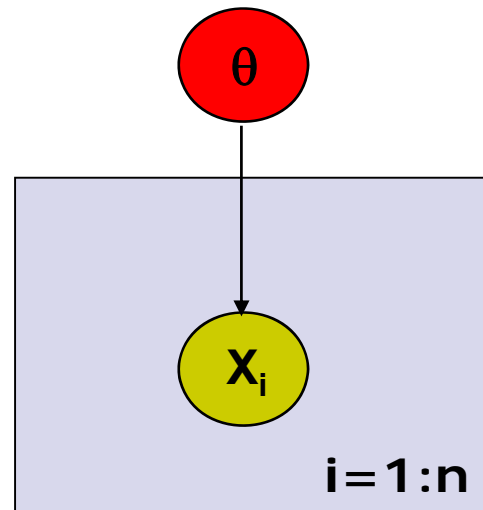
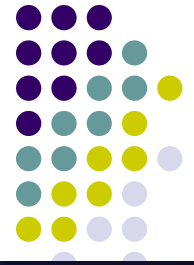
Conditionally Independent Observations



Model parameters

Data = $\{y_1, \dots, y_n\}$

“Plate” Notation



Model parameters

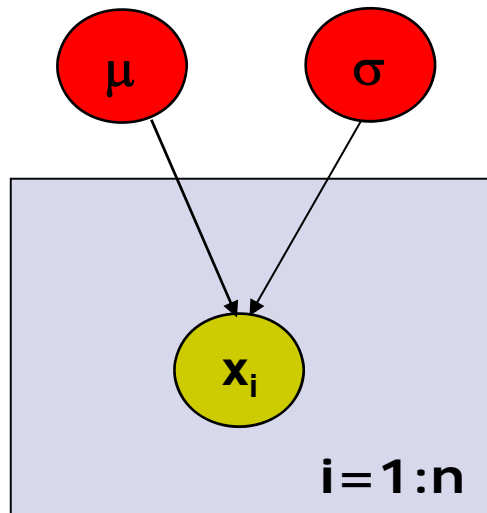
Data = $\{x_1, \dots, x_n\}$

Plate = rectangle in graphical model

**variables within a plate are replicated
in a conditionally independent manner**



Example: Gaussian Model



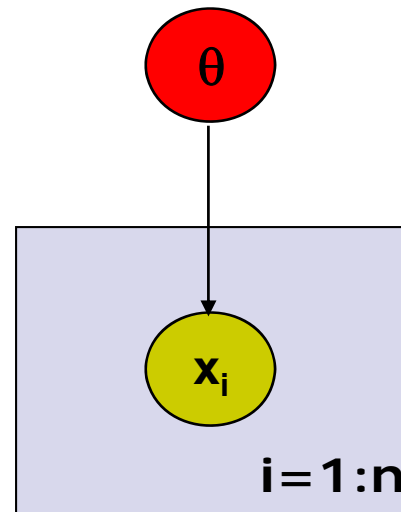
Generative model:

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

where $\theta = \{\mu, \sigma\}$

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

Bayesian models



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 GM 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
 - 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
- Support efficient inference and learning





Inference and Learning

- We now have compact representations of probability distributions: **BN**
- A BN M describes a unique probability distribution P
- Typical tasks:
 - Task 1: How do we answer **queries** about P ?
 - We use **inference** as a name for the process of computing answers to such queries
 - Task 2: How do we estimate a **plausible model** M from data D ?
 - i. We use **learning** as a name for the process of obtaining point estimate of M .
 - ii. But for *Bayesian*, they seek $p(M | D)$, which is actually an **inference** problem.
 - iii. When not all variables are observable, even computing point estimate of M need to do **inference** to impute the *missing data*.

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 $\mathbf{X} = \{X_1, X_2, \dots, X_n\}$
 - Without loss of generality $\mathbf{X}_v = \{X_{k+1}, \dots, X_n\}$,
 - Write $\mathbf{X}_H = \mathbf{X} \setminus \mathbf{X}_v$ as the set of hidden variables, \mathbf{X}_H can be \emptyset or \mathbf{X}

- Simplest query: compute probability of evidence

$$P(\mathbf{x}_v) = \sum_{\mathbf{X}_H} P(\mathbf{X}_H, \mathbf{x}_v) = \sum_{x_1} \dots \sum_{x_k} P(x_1, \dots, x_k, \mathbf{x}_v)$$

- this is often referred to as computing the **likelihood** of \mathbf{x}_v

Inferential Query 2: Conditional Probability



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

$$P(\mathbf{X}_H | \mathbf{X}_V = \mathbf{x}_V) = \frac{P(\mathbf{X}_H, \mathbf{x}_V)}{P(\mathbf{x}_V)} = \frac{P(\mathbf{X}_H, \mathbf{x}_V)}{\sum_{\mathbf{x}_H} P(\mathbf{X}_H = \mathbf{x}_H, \mathbf{x}_V)}$$

- this is the **a posteriori belief** in \mathbf{X}_H , given evidence \mathbf{x}_V
- We usually query a subset \mathbf{Y} of all hidden variables $\mathbf{X}_H = \{\mathbf{Y}, \mathbf{Z}\}$ and "don't care" about the remaining, \mathbf{Z} :

$$P(\mathbf{Y} | \mathbf{x}_V) = \sum_{\mathbf{z}} P(\mathbf{Y}, \mathbf{Z} = \mathbf{z} | \mathbf{x}_V)$$

- the process of summing out the "don't care" variables \mathbf{z} is called **marginalization**, and the resulting $P(\mathbf{Y} | \mathbf{x}_V)$ is called a **marginal prob.**

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 \mathbf{x}_V , and ignoring (the values of) other variables \mathbf{Z} :

$$Y^* | \mathbf{x}_V = \arg \max_y P(Y | \mathbf{x}_V) = \arg \max_y \sum_z P(Y, Z = z | \mathbf{x}_V)$$

- this is the **maximum a posteriori** configuration of Y .



Complexity of Inference

Thm:

Computing $P(X_H=x_H | x_v)$ in an arbitrary GM 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 GMs
 - For particular families of GMs, we can have provably efficient procedures



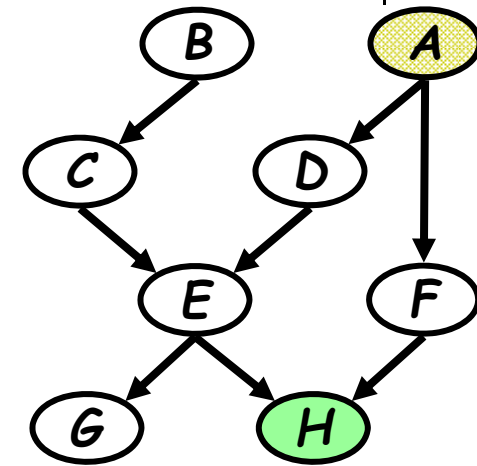
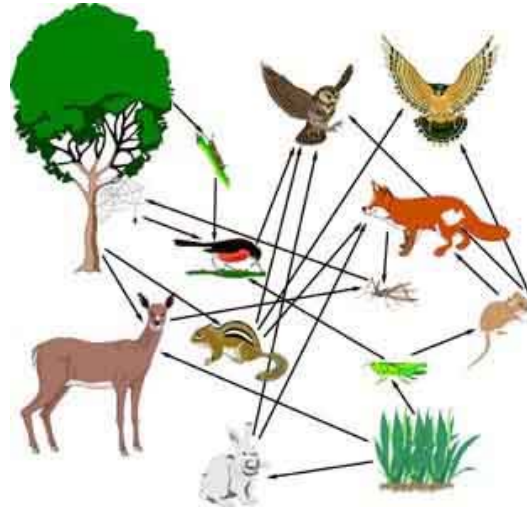
Approaches to inference

- Exact inference algorithms
 - The elimination algorithm
 - Belief propagation
 - The junction tree algorithms (but will not cover in detail here)
- Approximate inference techniques
 - Variational algorithms
 - Stochastic simulation / sampling methods
 - Markov chain Monte Carlo methods



Marginalization and Elimination

- A food web:



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

Query: $P(h)$

$$P(h) = \sum_g \sum_f \sum_e \sum_d \sum_c \sum_b \sum_a P(a, b, c, d, e, f, g, h)$$

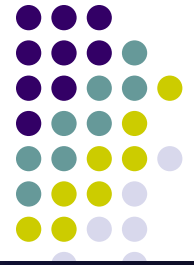


a naïve summation needs to enumerate over an exponential number of terms

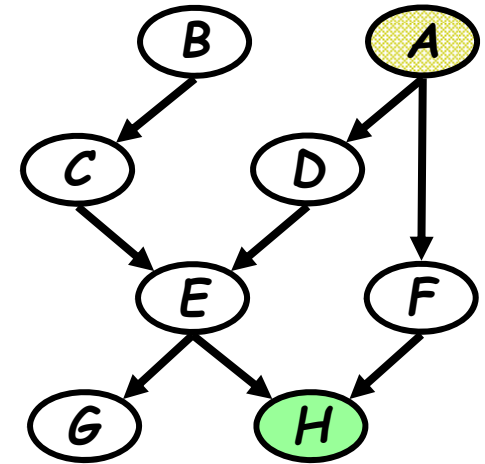
- By chain decomposition, we get

$$= \sum_g \sum_f \sum_e \sum_d \sum_c \sum_b \sum_a P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)P(h|e,f)$$

Variable Elimination



- Query: $P(A | h)$
 - Need to eliminate: B, C, D, E, F, G, H
- Initial factors:
 $P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)P(h|e,f)$
- Choose an elimination order: H, G, F, E, D, C, B

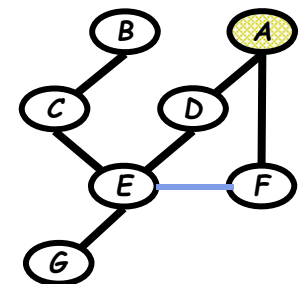


- Step 1:
 - **Conditioning** (fix the evidence node (i.e., h) on its observed value (i.e., \tilde{h})):

$$m_h(e, f) = p(h = \tilde{h} | e, f)$$

- This step is isomorphic to a marginalization step:

$$m_h(e, f) = \sum_h p(h | e, f) \delta(h = \tilde{h})$$

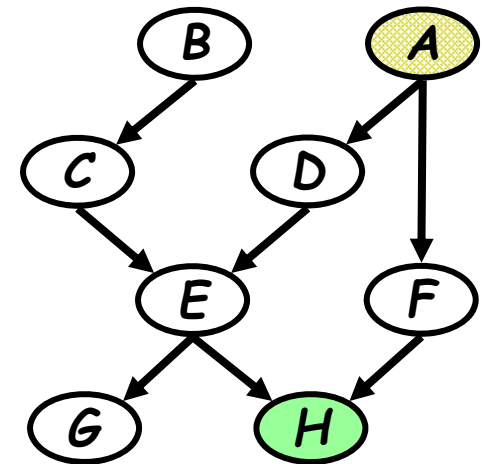




Example: Variable Elimination

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

$$P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f)$$
$$\Rightarrow P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)\underline{m_h(e, f)}$$

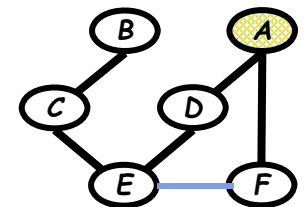


- Step 2: Eliminate G

- compute

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

$$\Rightarrow P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)m_g(e)m_h(e, f)$$
$$= P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)\underline{m_h(e, f)}$$



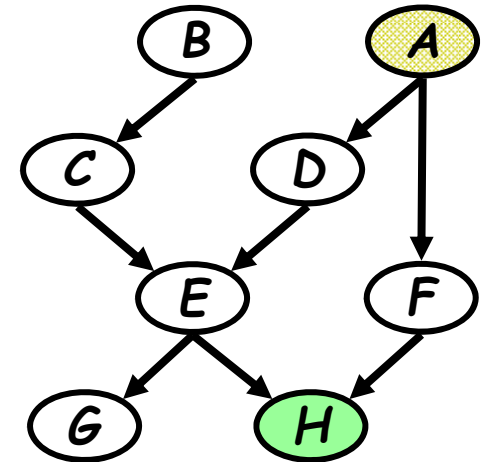


Example: Variable Elimination

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

- Initial factors:

$$\begin{aligned} &P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f) \\ \Rightarrow &P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)\underline{m_h(e, f)} \end{aligned}$$

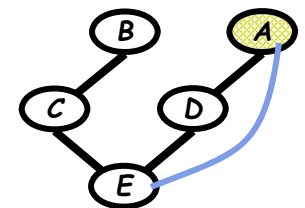


- Step 3: Eliminate F

- compute

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

$$\Rightarrow P(a)P(b)P(c | b)P(d | a)P(e | c, d)\underline{m_f(a, e)}$$





Example: Variable Elimination

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

- Initial factors:

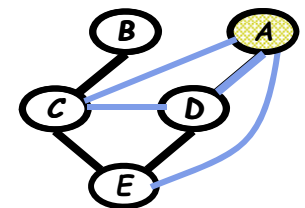
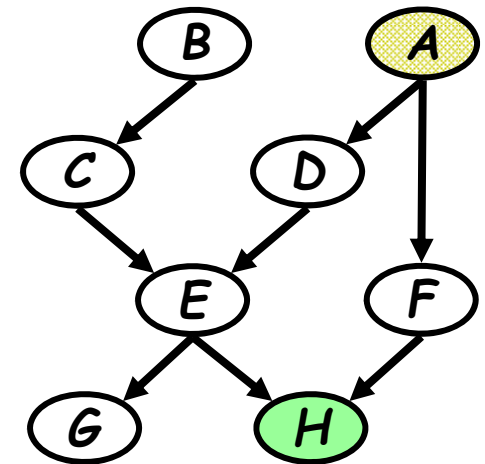
$$\begin{aligned} &P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f) \\ \Rightarrow &P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)P(g | e)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | b)P(d | a)P(e | c, d)P(f | a)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | b)P(d | a)\underline{P(e | c, d)m_f(a, e)} \end{aligned}$$

- Step 4: Eliminate E

- compute

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

$$\Rightarrow P(a)P(b)P(c | b)P(d | a)\underline{m_e(a, c, d)}$$



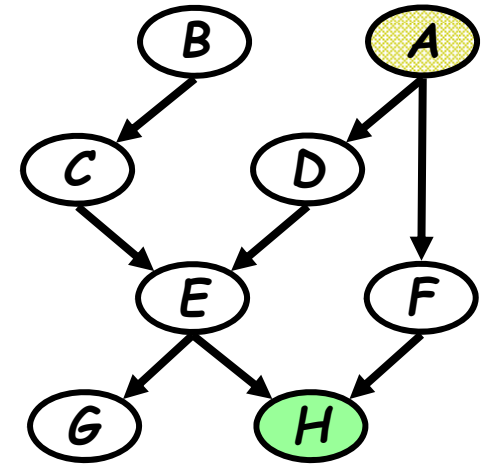


Example: Variable Elimination

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

- Initial factors:

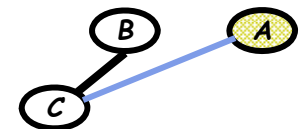
$$\begin{aligned} &P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)P(h|e,f) \\ \Rightarrow &P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)P(g|e)m_h(e,f) \\ \Rightarrow &P(a)P(b)P(c|b)P(d|a)P(e|c,d)P(f|a)m_h(e,f) \\ \Rightarrow &P(a)P(b)P(c|b)P(d|a)P(e|c,d)m_f(a,e) \\ \Rightarrow &P(a)P(b)P(c|b)P(d|a)m_e(a,c,d) \end{aligned}$$

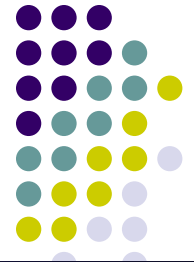


- Step 5: Eliminate D

- compute
$$m_d(a,c) = \sum_d p(d|a)m_e(a,c,d)$$

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





Example: Variable Elimination

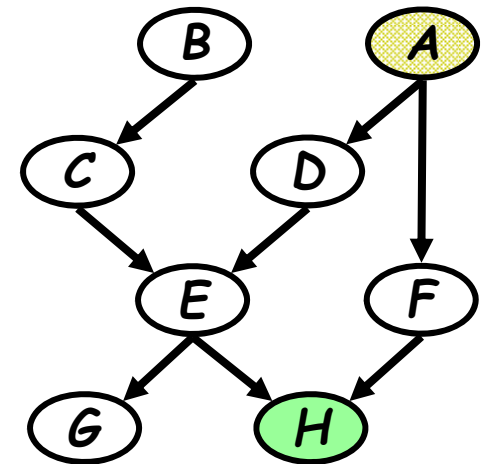
- Query: $P(B | h)$
 - Need to eliminate: B, C
- Initial factors:

$$\begin{aligned} &P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)P(g | e)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)P(e | c, d)m_f(a, e) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)m_e(a, c, d) \\ \Rightarrow &P(a)P(b)P(c | d)m_d(a, c) \end{aligned}$$

- Step 6: **Eliminate C**

- compute $m_c(a, b) = \sum_c p(c | b)m_d(a, c)$

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





Example: Variable Elimination

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

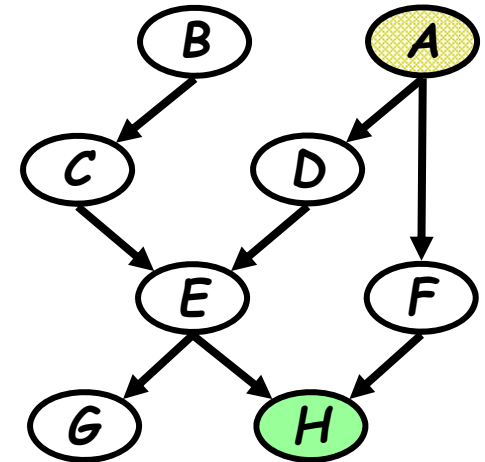
$$\begin{aligned} & P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f) \\ \Rightarrow & P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)P(g | e)m_h(e, f) \\ \Rightarrow & P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)m_h(e, f) \\ \Rightarrow & P(a)P(b)P(c | d)P(d | a)P(e | c, d)m_f(a, e) \\ \Rightarrow & P(a)P(b)P(c | d)P(d | a)m_e(a, c, d) \\ \Rightarrow & P(a)P(b)P(c | d)m_d(a, c) \\ \Rightarrow & P(a)P(b)m_c(a, b) \end{aligned}$$

- Step 7: **Eliminate B**

- compute

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

$$\Rightarrow \underline{P(a)m_b(a)}$$



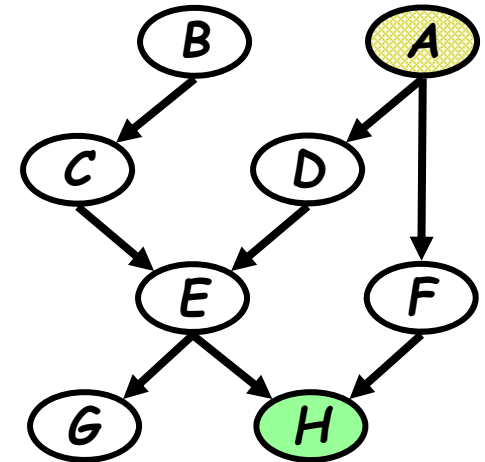


Example: Variable Elimination

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

$$\begin{aligned} &P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)P(g | e)P(h | e, f) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)P(g | e)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)P(e | c, d)P(f | a)m_h(e, f) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)P(e | c, d)m_f(a, e) \\ \Rightarrow &P(a)P(b)P(c | d)P(d | a)m_e(a, c, d) \\ \Rightarrow &P(a)P(b)P(c | d)m_d(a, c) \\ \Rightarrow &P(a)P(b)m_c(a, b) \\ \Rightarrow &P(a)m_b(a) \end{aligned}$$

- Step 8: **Wrap-up** $p(a, \tilde{h}) = p(a)m_b(a), \quad p(\tilde{h}) = \sum_a p(a)m_b(a)$
 $\Rightarrow P(a | \tilde{h}) = \frac{p(a)m_b(a)}{\sum_a p(a)m_b(a)}$



Complexity of variable elimination



- Suppose in one elimination step we compute

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

This requires

- $k \cdot |\text{Val}(X)| \cdot \prod_i |\text{Val}(\mathbf{y}_{C_i})|$ **multiplications**
 - For each value of x, y_1, \dots, y_k , we do k multiplications
- $|\text{Val}(X)| \cdot \prod_i |\text{Val}(\mathbf{y}_{C_i})|$ **additions**
 - For each value of y_1, \dots, y_k , we do $|\text{Val}(X)|$ additions

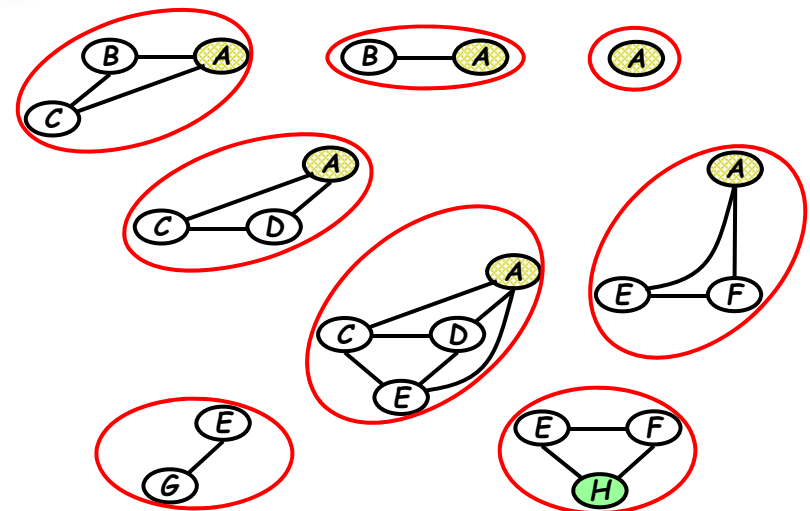
Complexity is **exponential** in number of variables in the intermediate factor



Elimination Clique

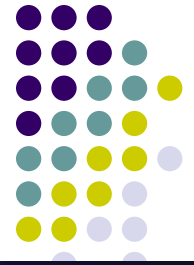
- Induced dependency during marginalization is captured in elimination cliques
 - Summation \leftrightarrow elimination
 - Intermediate term \leftrightarrow elimination clique

$$\begin{aligned} & P(a)P(b)P(c|b)P(d|a)P(e|c, d)P(f|a)P(g|e)P(h|e, f) \\ \Rightarrow & P(a)P(b)P(c|b)P(d|a)P(e|c, d)P(f|a)P(g|e)\phi_h(e, f) \\ \Rightarrow & P(a)P(b)P(c|b)P(d|a)P(e|c, d)P(f|a)\phi_g(e)\phi_h(e, f) \\ \Rightarrow & P(a)P(b)P(c|b)P(d|a)P(e|c, d)\phi_f(a, e) \\ \Rightarrow & P(a)P(b)P(c|b)P(d|a)\phi_e(a, c, d) \\ \Rightarrow & P(a)P(b)P(c|b)\phi_d(a, c) \\ \Rightarrow & P(a)P(b)\phi_c(a, b) \\ \Rightarrow & P(a)\phi_b(a) \\ \Rightarrow & \phi(a) \end{aligned}$$



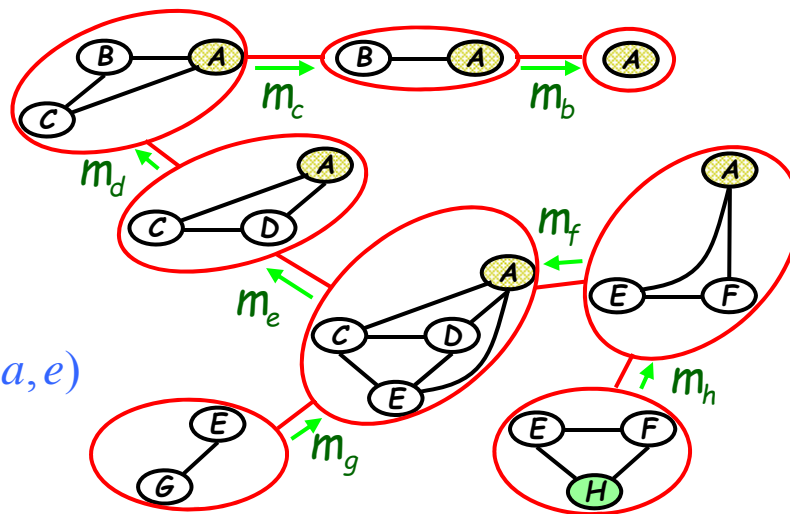
- Can this lead to a generic inference algorithm?

From Elimination to Message Passing



- Elimination \equiv message passing on a **clique tree**

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

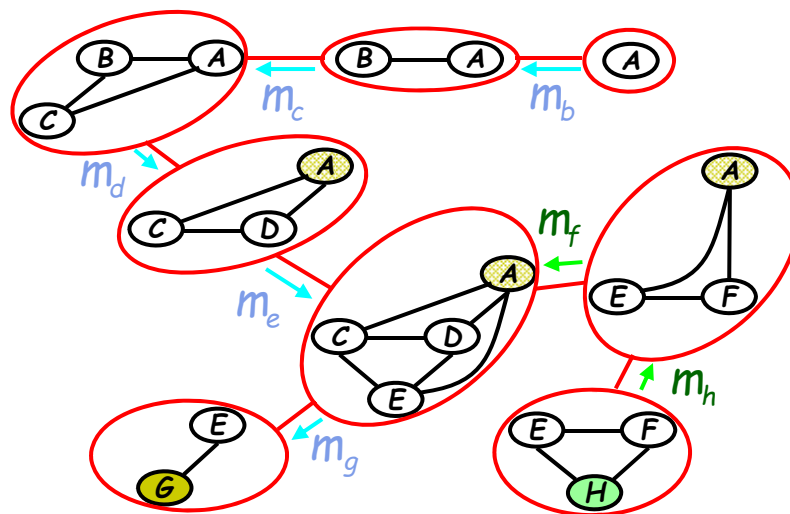


- Messages can be reused

From Elimination to Message Passing

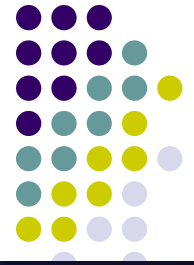


- Elimination \equiv message passing on a **clique tree**
 - Another query ...



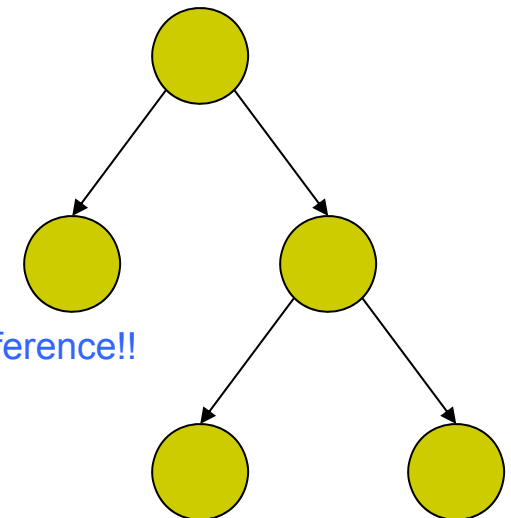
- Messages m_f and m_h are reused, others need to be recomputed

From elimination to message passing

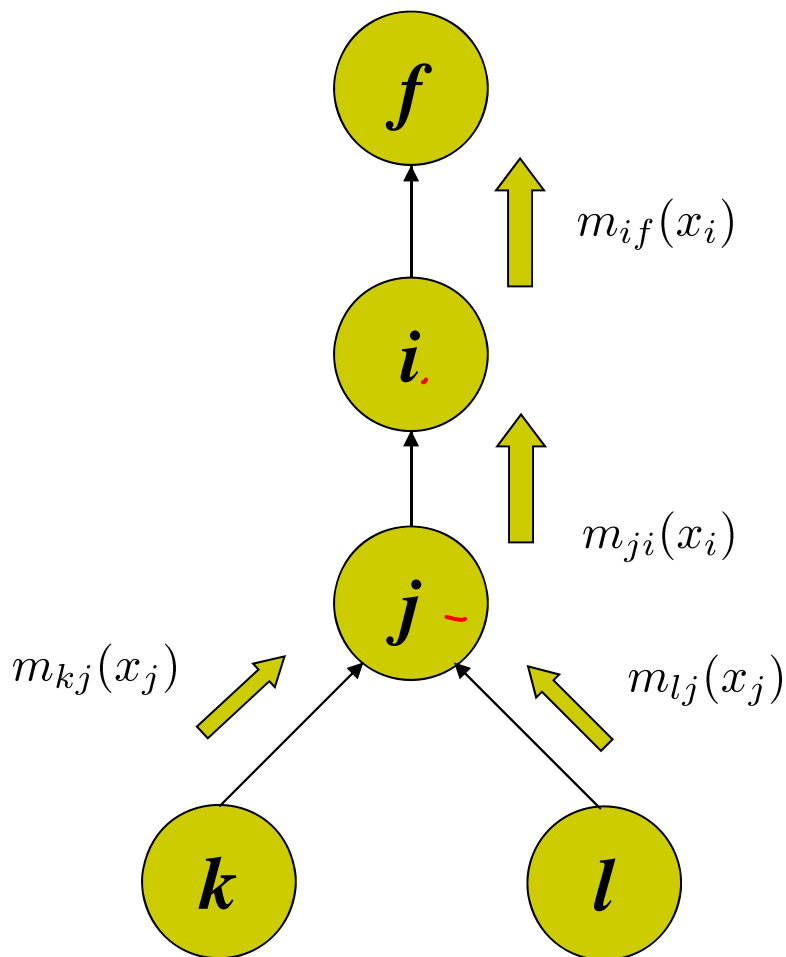


- Recall **ELIMINATION** algorithm:
 - Choose an ordering \mathcal{Z} in which query node f is the final node
 - Place all potentials on an active list
 - Eliminate node i by removing all potentials containing i , take sum/product over x_i .
 - Place the resultant factor back on the list

 - For a **TREE** graph:
 - Choose query node f as the root of the tree
 - View tree as a directed tree with edges pointing towards from f
 - Elimination ordering based on depth-first traversal
 - Elimination of each node can be considered as **message-passing (or Belief Propagation)** directly along tree branches, rather than on some transformed graphs
- thus, we can use the tree itself as a data-structure to do general inference!!



Message passing for trees



Let $m_{ij}(x_i)$ denote the factor resulting from eliminating variables from below up to i , which is a function of x_i :

$$m_{ji}(x_i) = \sum_{x_j} \left(\psi(x_j) \psi(x_i, x_j) \prod_{k \in N(j) \setminus i} m_{kj}(x_j) \right)$$

This is reminiscent of a **message** sent from j to i .

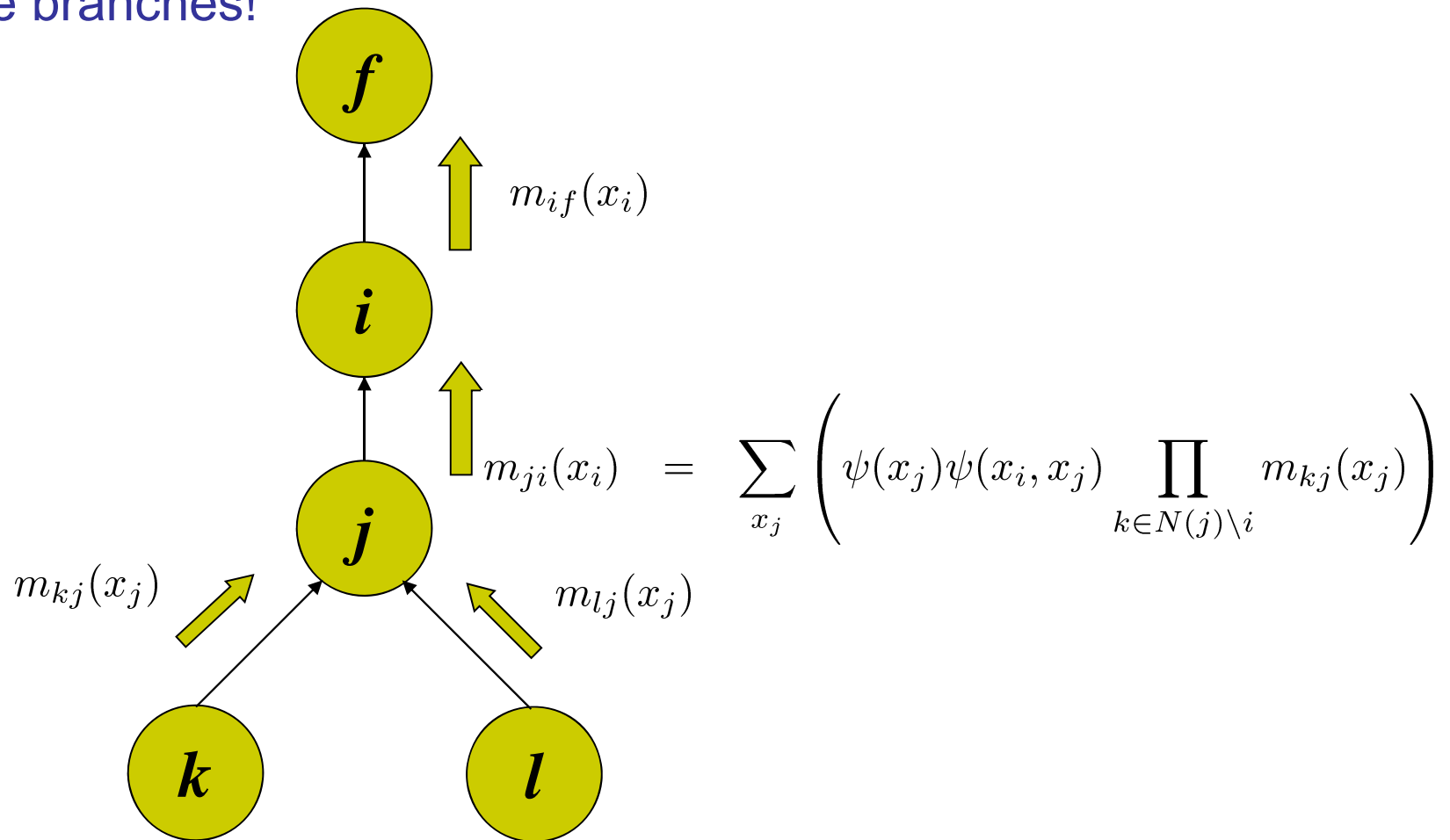
$$m_{ji}(x_i) = \sum_{x_j} \left(\psi(x_j) \psi(x_i, x_j) \prod_{k \in N(j) \setminus i} m_{kj}(x_j) \right)$$

$$p(x_f) \propto \psi(x_f) \prod_{e \in N(f)} m_{ef}(x_f)$$

$m_{ij}(x_i)$ represents a "belief" of x_i from x_j !



- Elimination on trees is equivalent to message passing along tree branches!

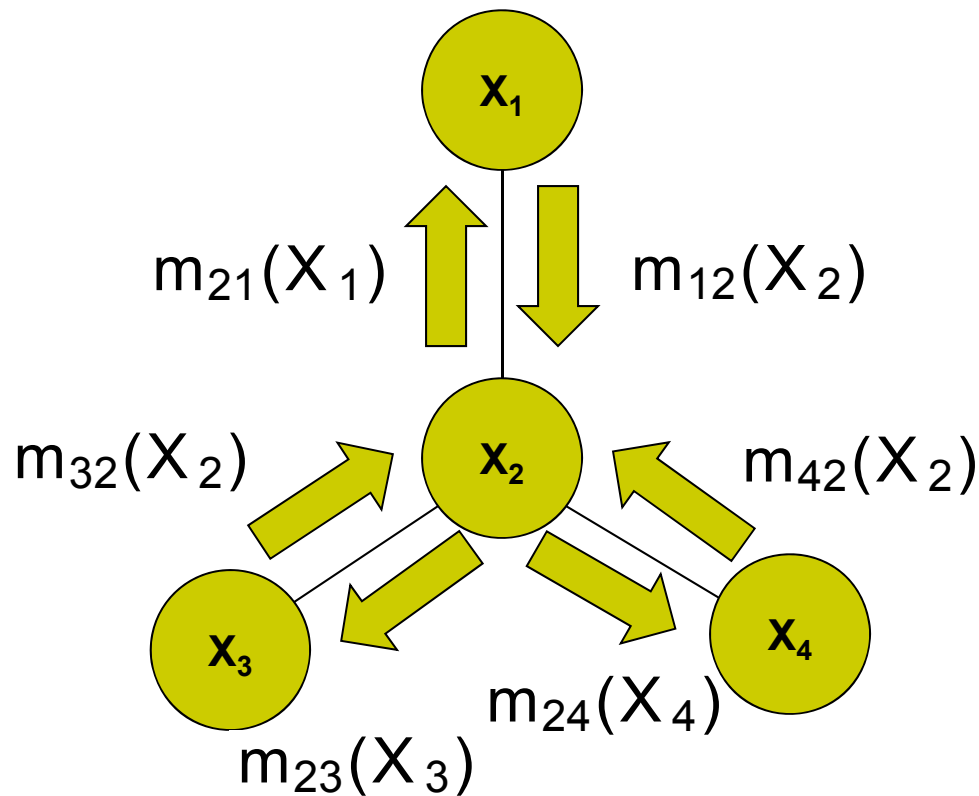


$$= \sum_{x_j} \left(\psi(x_j) \psi(x_i, x_j) \prod_{k \in N(j) \setminus i} m_{kj}(x_j) \right)$$

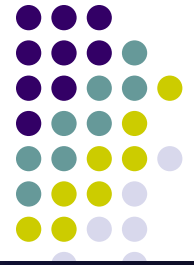


The message passing protocol:

- A two-pass algorithm:



Belief Propagation (SP-algorithm): Sequential implementation



```

SUM-PRODUCT( $\mathcal{T}, E$ )
  EVIDENCE( $E$ )
   $f = \text{CHOOSEROOT}(\mathcal{V})$ 
  for  $e \in \mathcal{N}(f)$ 
    COLLECT( $f, e$ )
  for  $e \in \mathcal{N}(f)$ 
    DISTRIBUTE( $f, e$ )
  for  $i \in \mathcal{V}$ 
    COMPUTEMARGINAL( $i$ )
  
```

```

EVIDENCE( $E$ )
  for  $i \in E$ 
     $\psi^E(x_i) = \psi(x_i)\delta(x_i, \bar{x}_i)$ 
  for  $i \notin E$ 
     $\psi^E(x_i) = \psi(x_i)$ 
  
```

```

COLLECT( $i, j$ )
  for  $k \in \mathcal{N}(j) \setminus i$ 
    COLLECT( $j, k$ )
  SENDMESSAGE( $j, i$ )
  
```

```

DISTRIBUTE( $i, j$ )
  SENDMESSAGE( $i, j$ )
  for  $k \in \mathcal{N}(j) \setminus i$ 
    DISTRIBUTE( $j, k$ )
  
```

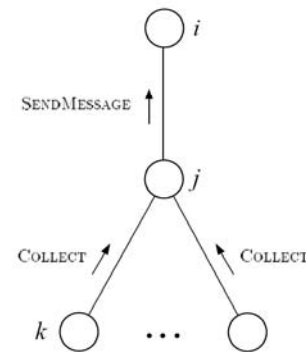
```

SENDMESSAGE( $j, i$ )
 $m_{ji}(x_i) = \sum_{x_j} (\psi^E(x_j)\psi(x_i, x_j)) \prod_{k \in \mathcal{N}(j) \setminus i} m_{kj}(x_j)$ 
  
```

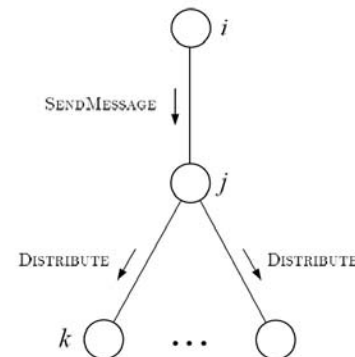
```

COMPUTEMARGINAL( $i$ )
 $p(x_i) \propto \psi^E(x_i) \prod_{j \in \mathcal{N}(i)} m_{ji}(x_i)$ 
  
```

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Inference on general GM

- Now, what if the GM is not a tree-like graph?
- Can we still directly run message message-passing protocol along its edges?
- For non-trees, we do not have the guarantee that message-passing will be consistent!
- Then what?
 - Construct a graph data-structure from P that has a tree structure, and run message-passing on it!

→ Junction tree algorithm

Summary



- The simple Eliminate algorithm captures the key algorithmic Operation underlying probabilistic inference:
--- That of taking a sum over product of potential functions
- The computational complexity of the Eliminate algorithm can be reduced to purely graph-theoretic considerations.
- This graph interpretation will also provide hints about how to design improved inference algorithms
- What can we say about the overall computational complexity of the algorithm? In particular, how can we control the "size" of the summands that appear in the sequence of summation operation.