

GRAPHICAL MODELS

- Bayesian networks
 - Generative models
 - Discriminative models
 - Markov chains
 - D-separation
 - Hidden Markov Models

- Undirected models
 - Factorization
 - Hammersley-Clifford Theorem
 - Moralization

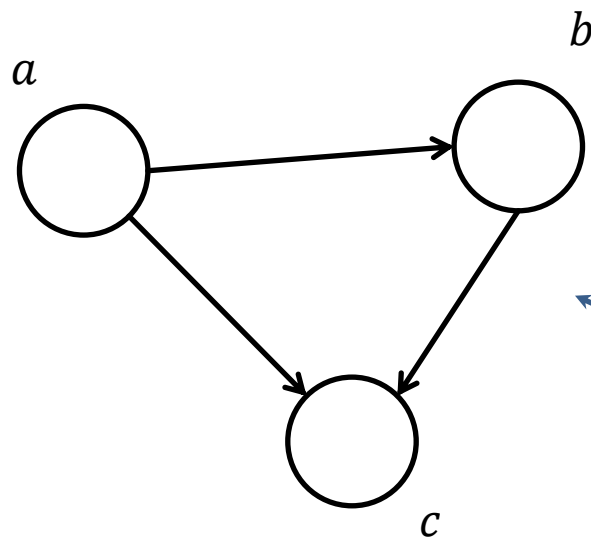
- What is a graphical model?
 - Graph $G(V, E)$
 - V : Set of random variables
 - $E(\subseteq V \times V)$: Set of dependence relationships

- Why graphical models?
 - Visualize the structure of probabilistic models
 - Obtain insights into properties of variables and their interdependencies (e.g. **conditional independence, causality**) through **graph-theoretic** means
 - Perform **probabilistic inference** through graphical manipulations

- Types of graphical models
 - **Directed** graphical models \equiv **Bayesian networks** (coined by Judea Pearl 1985)
 - **Undirected** graphical models \equiv **Markov random fields**

Bayesian networks (1)

- Represent conditional dependencies between random variables



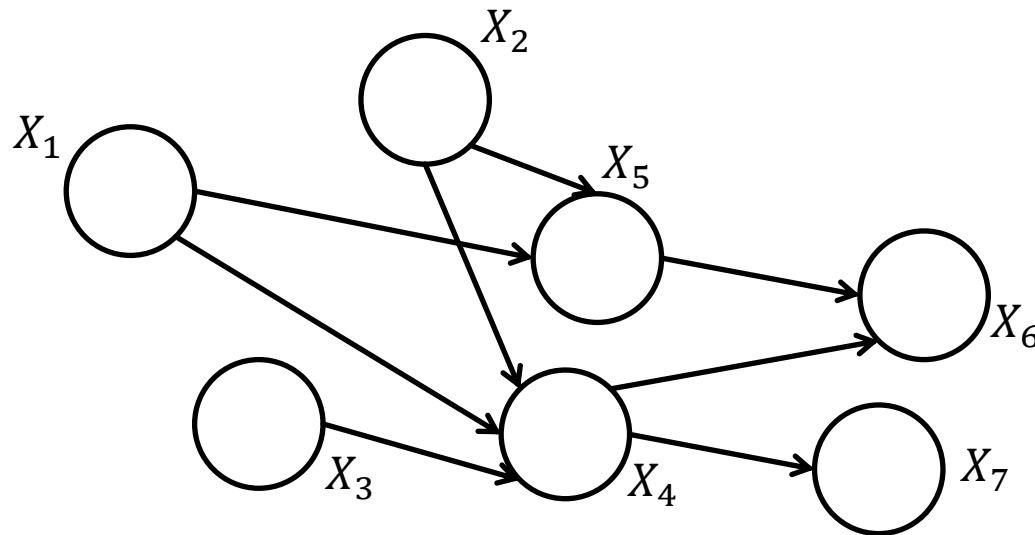
Example from
C. Bishop: PRML book

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

$$P(X_1, \dots, X_k) = P(X_k|X_1, \dots, X_{k-1}) \dots P(X_2|X_1)P(X_1)$$

Bayesian networks (2)

- Represent causal dependencies between random variables
- **Local Markov Property:** Each variable is conditionally independent of its non-descendants given its parents

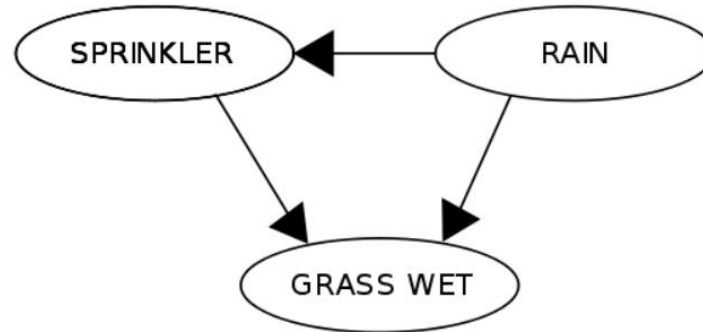


$$P(X_1, \dots, X_7) = P(X_1)P(X_2)P(X_3)P(X_4|X_1, X_2, X_3)P(X_5|X_1, X_2)P(X_6|X_4, X_5)P(X_7|X_4)$$

➔ Generally: $P(\mathbf{X}) = \prod_{i=1}^k P(X_i | \text{par}_i)$, where par_i denote the parents of X_i

Bayesian network example

RAIN	SPRINKLER	
	T	F
F	0.4	0.6
T	0.01	0.99



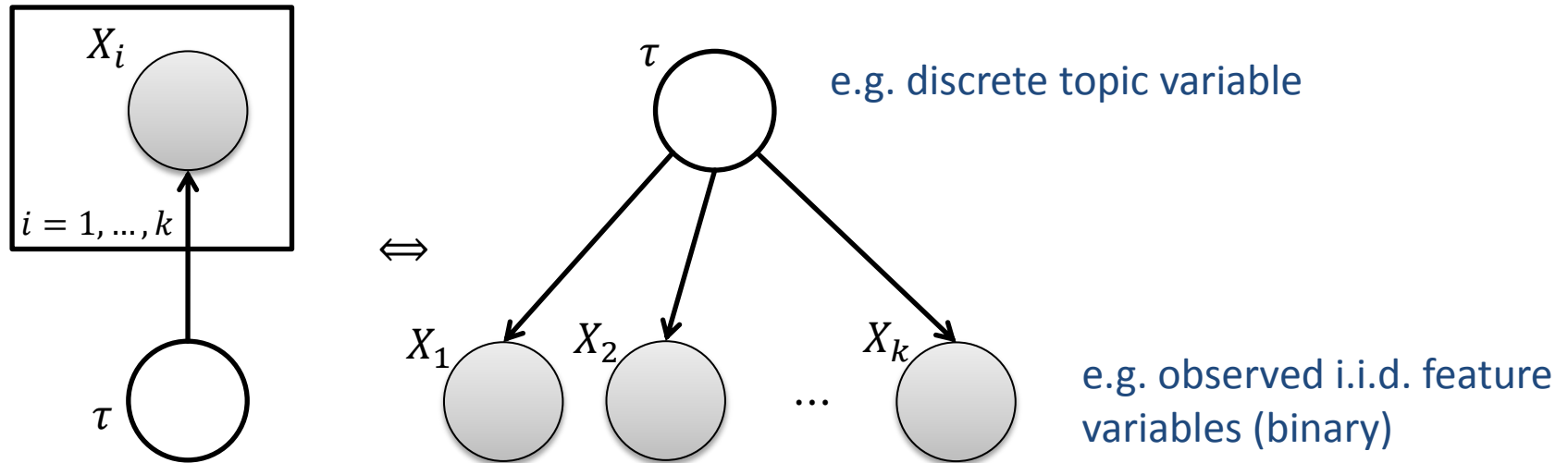
RAIN	
T	F
0.2	0.8

		GRASS WET	
SPRINKLER	RAIN	T	F
F	F	0.0	1.0
F	T	0.8	0.2
T	F	0.9	0.1
T	T	0.99	0.01

$$P(S, R, G) = P(R)P(S|R)P(G|S, R)$$

- What is the probability that the sprinkler was on given that the grass is wet?

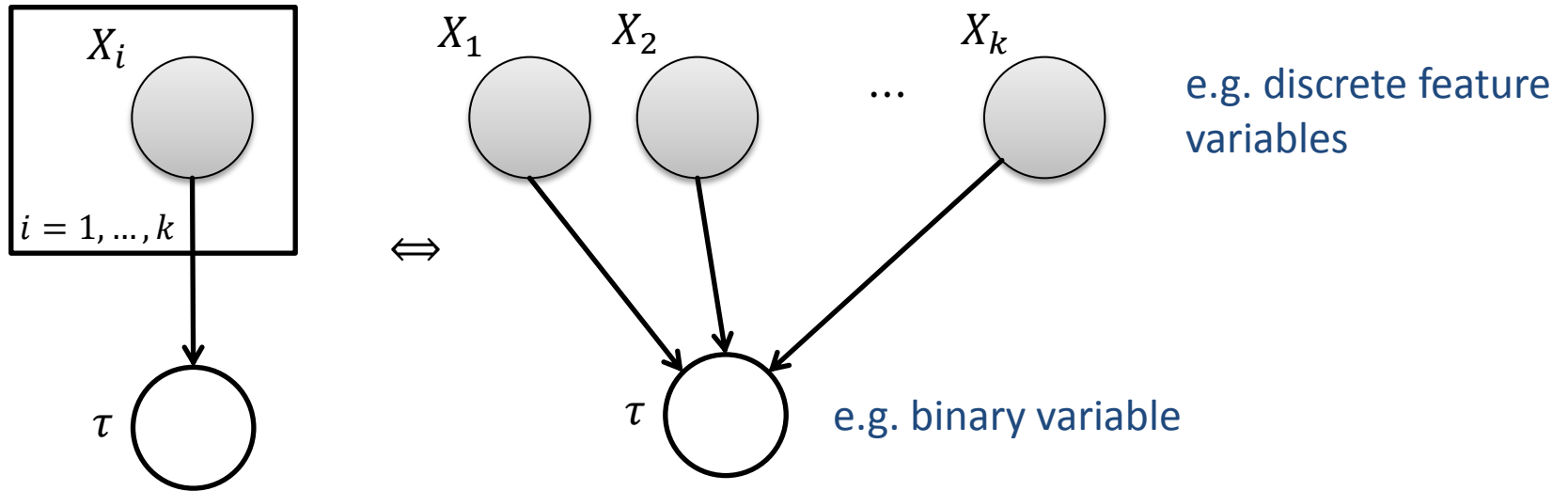
Generative models



$$P(X_1, \dots, X_k, \tau) = P(\tau) \prod_{i=1}^k P(X_i | \tau)$$

... we need to estimate $k + 1$ parameters

Discriminative models



$$P(\tau = 1 | X_1, \dots, X_k)$$

... we need to estimate l^k parameters, if each X_i has l states

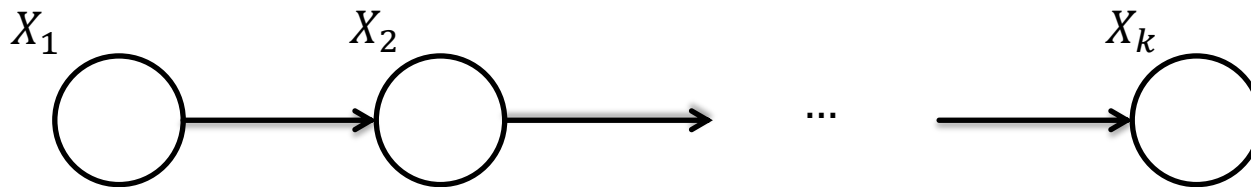
The trick is to parameterize each X_i with a weight β_i and estimate

$$\frac{1}{1 + \exp(-\beta_0 - \beta_1 X_1 - \dots - \beta_k X_k)}$$

... we need to estimate $k + 1$ parameters

Markov chains

- Generally, for the joint distribution of k discrete variables X_1, \dots, X_k , with l states each, we need to estimate $l^k - 1$ parameters
- For k -node Markov chain: $l - 1 + (k - 1)l(l - 1)$ parameters



- A Markov chain is a Bayesian network in which each variable has at most one predecessor

Conditional independence

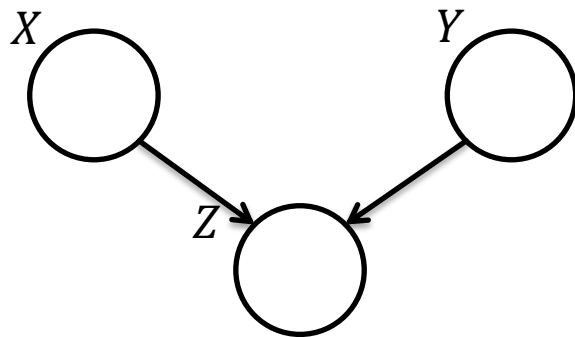
X is independent of Y given Z (we write: $X \perp Y|Z$)

\Leftrightarrow

$$P(X|Y, Z) = P(X|Z)$$

\Leftrightarrow

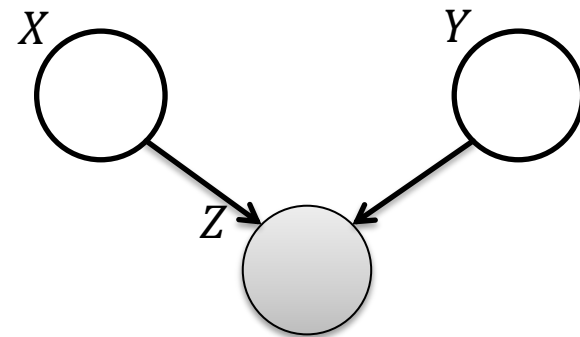
$$P(X, Y|Z) = P(X|Z)P(Y|Z)$$



$$P(X, Y, Z) = P(X)P(Y)P(Z|X, Y)$$

\Leftrightarrow

$$P(X, Y) = P(X)P(Y) \Leftrightarrow X \perp Y|\emptyset$$



$$P(X, Y|Z) = P(X, Y, Z)/P(Z)$$

\Leftrightarrow

$$P(X, Y|Z) = P(X)P(Y)P(Z|X, Y)/P(Z) \Rightarrow X \not\perp Y|Z$$

Conditional independence example

➤ Example from C. Bishop, PRML book

B : battery (0=flat, 1=full)

F : fuel (0=empty, 1=full)

R : Fuel reading (0=empty, 1=full)

$P(B = 1) = 0.9$

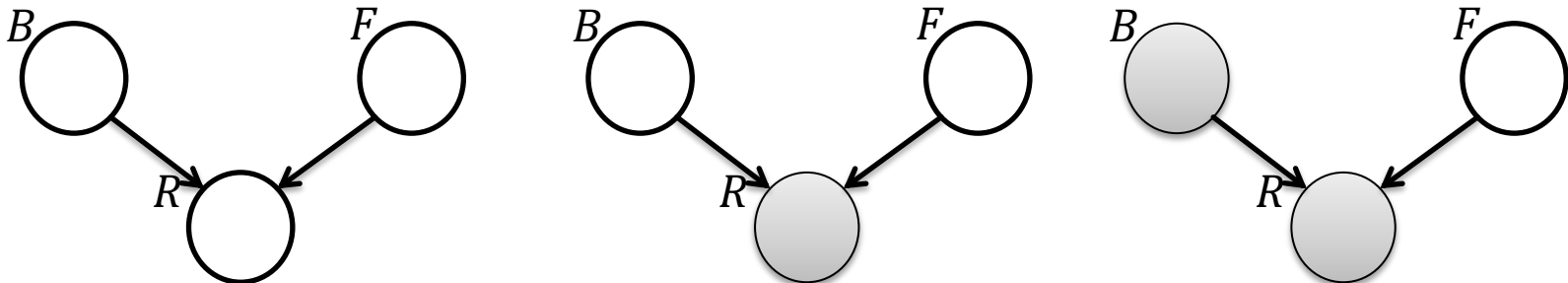
$P(F = 1) = 0.9$

$P(R = 1|B = 1, F = 1) = 0.8$

$P(R = 1|B = 1, F = 0) = 0.2$

$P(R = 1|B = 0, F = 1) = 0.2$

$P(R = 1|B = 0, F = 0) = 0.1$



We can calculate:

$$P(R = 0) = \sum_{B \in \{0,1\}} \sum_{F \in \{0,1\}} P(R = 0|B, F)P(B)P(F) = 0.315$$

$$P(R = 0|F = 0) = \sum_{B \in \{0,1\}} P(R = 0|B, F = 0)P(B) = 0.81$$

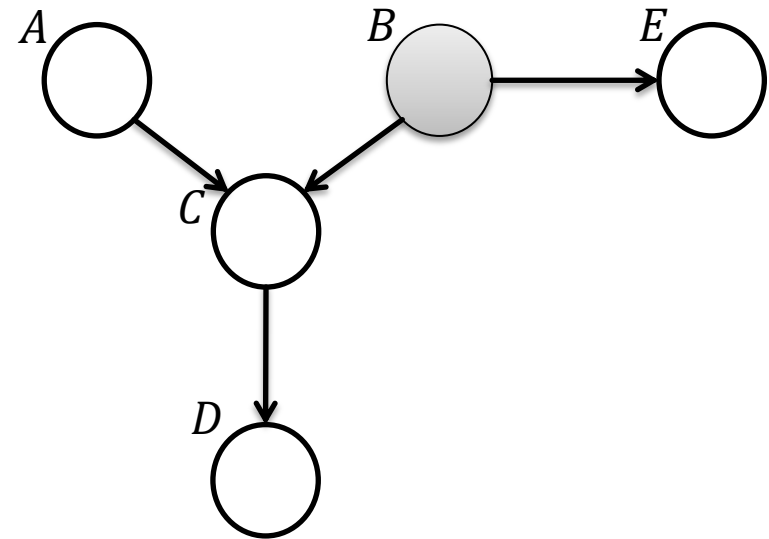
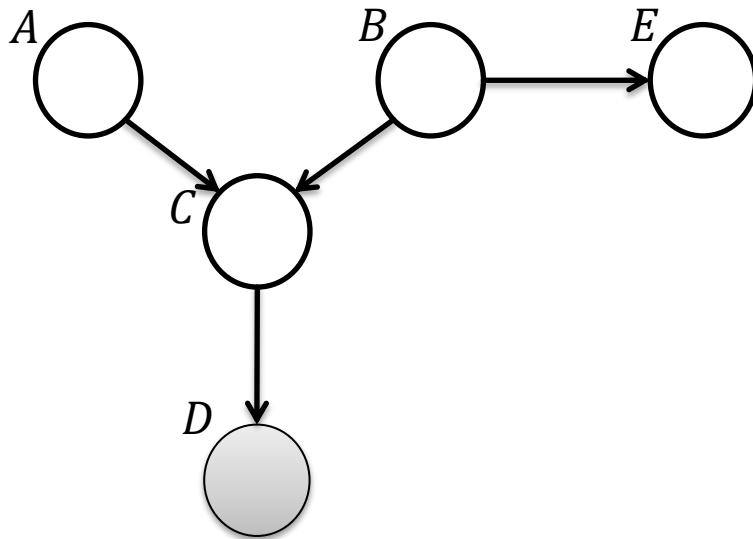
$$P(F = 0|R = 0) = \frac{P(R=0|F=0)P(F=0)}{P(R=0)} \approx 0.257 \text{ (probability of } F = 0 \text{ increases)}$$

$$P(F = 0|R = 0, B = 0) \approx 0.111 \text{ (prob. of } F = 0 \text{ decreases, is "explained away")}$$

D-separation

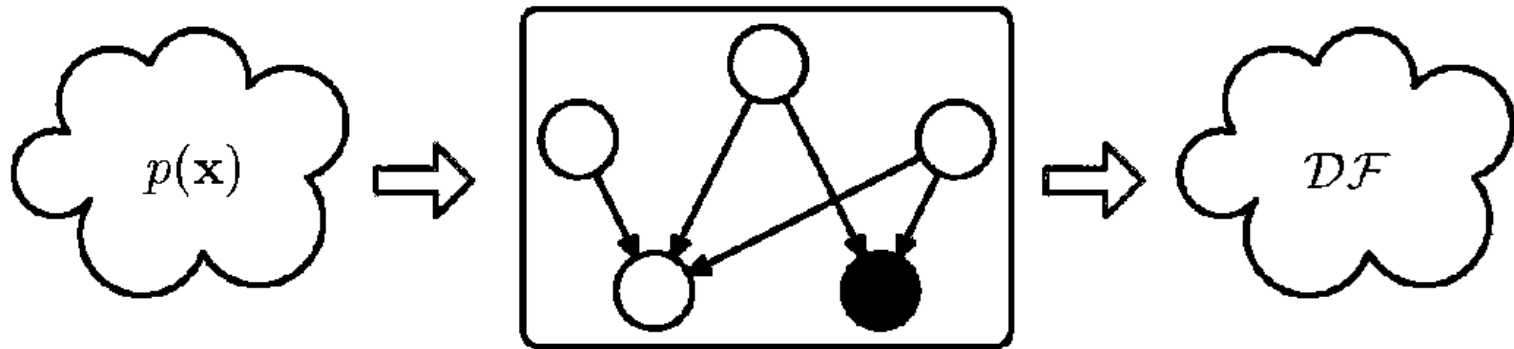
- Let A, B, C be disjoint subsets of nodes from a Bayesian network
- A path from A to B is **blocked** if it contains node v such that either
 - arrows on the path meet head-to-tail ($\rightarrow v \rightarrow$) or tail-to tail ($\leftarrow v \rightarrow$) at v and v is from C , or
 - arrows on the path meet head-to-head at v ($\rightarrow v \leftarrow$) and neither v nor any of its descendants are in C
- A is **d-separated** from B by C if all paths from A to B are **blocked**
- **Theorem:** If A is d-separated from B by C then $A \perp B | C$

D-separation examples



- $A \perp E | D$ ❌
- $A \perp E | C$ ❌
- $A \perp E | B$ ✅

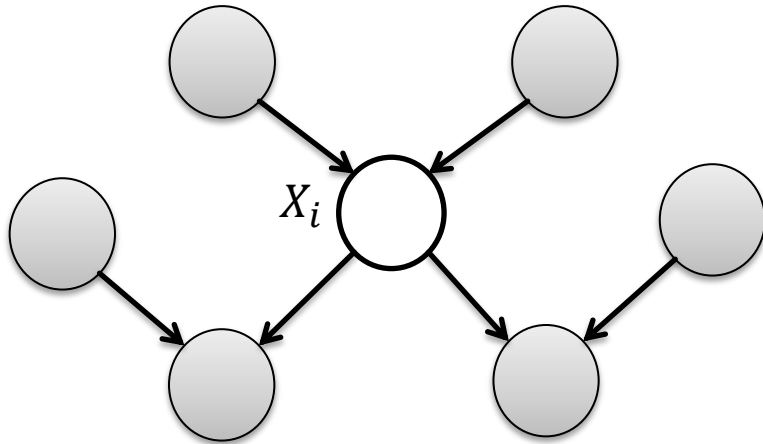
Bayesian networks as distribution filters



From C. Bishop: PRML book

- Any joint probability distribution $P(\mathbf{x})$ that factorizes according to the graphical model passes through the filter
- The set of distributions $P(\mathbf{x})$ that pass through the filter is denoted by \mathcal{DF} ; this is exactly the set of distributions that respect all conditional independencies implied by the d-separation

Markov blanket



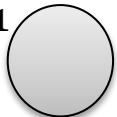
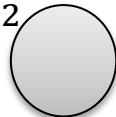
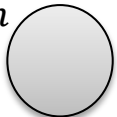
$$P(X_i | \{X_{j \neq i}\}) = \frac{P(X_1, \dots, X_k)}{\int P(X_1, \dots, X_k) dX_i}$$
$$= \frac{\prod_j P(X_j | \text{par}_j)}{\int \prod_j P(X_j | \text{par}_j) dX_i}$$

Factors independent of X_i cancel out!

- In a Bayesian network, a variable is independent of all other variables given its Markov blanket

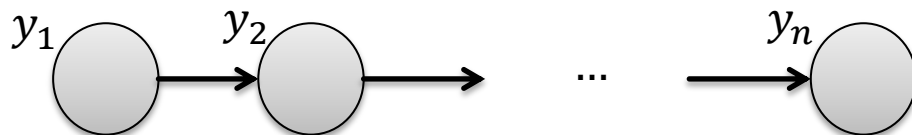
Hidden Markov models (HMMs)

- Useful for explaining sequential data, e.g., arising from
 - Measurements of time series
 - Observations of sequential nucleotide base pairs along a DNA strand
 - Sequential tokens in a sentence/speech
- Simplest way to model the probability of an observed sequence y_1, \dots, y_n is to assume pairwise independence between the observations:

y_1  y_2  ... y_n  $P(\mathbf{y}) = \prod_{i=1}^n P(y_i)$

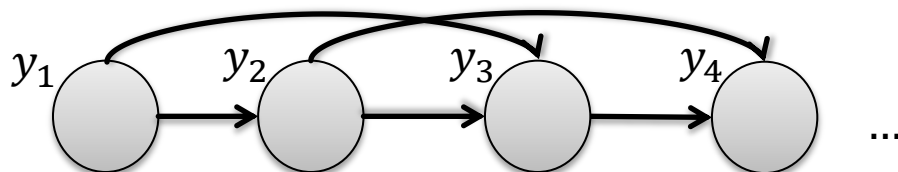
- Above assumption may be too strong; if it is relaxed such that every observation is dependent only on the (most) recent observations we obtain a **Markov model**

From Markov models to HMMs



First-order Markov chain

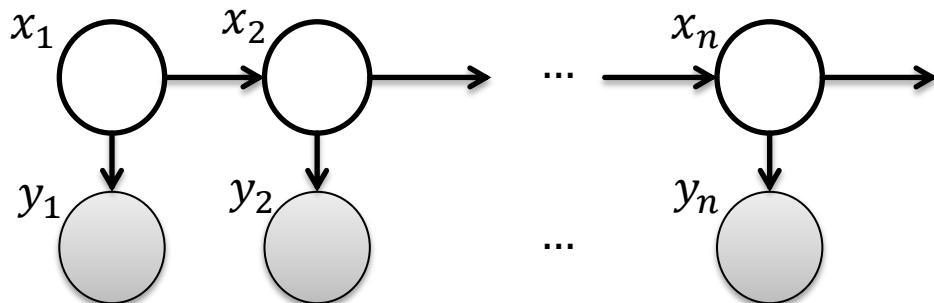
$$P(\mathbf{y}) = P(y_1) \prod_{i=2}^n P(y_i | y_{i-1})$$



Second-order Markov chain

$$P(\mathbf{y}) = P(y_1)P(y_2 | y_1) \prod_{i=3}^n P(y_i | y_{i-1}, y_{i-2})$$

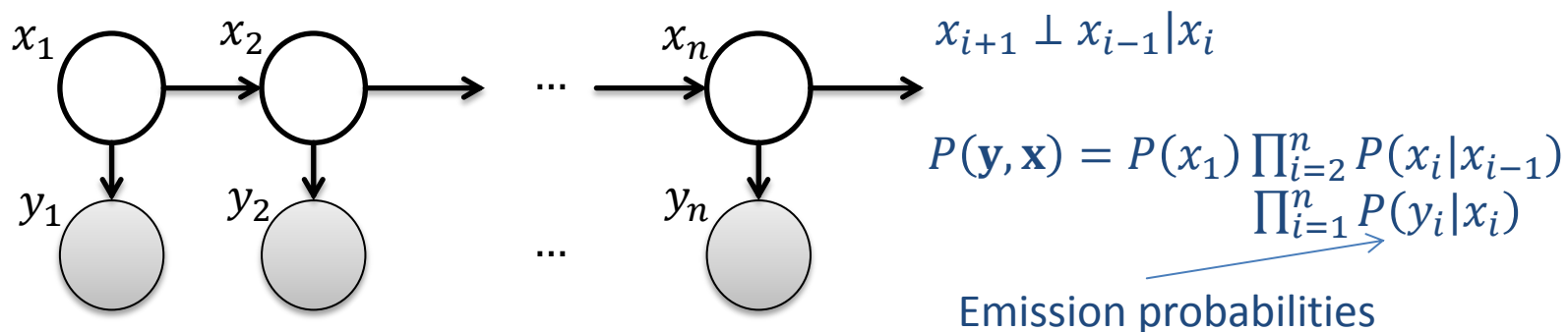
- More powerful model assumes latent variables (unknown states) for each y_i :



$$P(\mathbf{y}, \mathbf{x}) = P(x_1) \prod_{i=2}^n P(x_i | x_{i-1}) \prod_{i=1}^n P(y_i | x_i)$$

Extending Markov chains to HMMs

- If the latent variables are discrete the above model is an HMM



➤ Inference

- 1) Find the most likely state at a given x_i : $\max_s P(x_i = s | y_1, \dots, y_i)$
(can be solved with the forward-backward algorithm)
- 2) Find the most likely sequence of states: $\max_s P(\mathbf{x} = \mathbf{s}, y_1, \dots, y_i)$
(can be solved with the Viterbi algorithm)

The Viterbi (dynamic programming) algorithm

- Input: observation space $O = \{o_1, \dots, o_N\}$, observations $Y = \{y_1, \dots, y_n\}$, state space $S = \{s_1, \dots, s_k\}$, $k \times k$ -dim. state transition matrix \mathbf{A} , $k \times N$ -dim. emission matrix \mathbf{B} , two $k \times n$ -dim. matrices T_1, T_2 , array $\boldsymbol{\pi}$ of size k (with prior probabilities for each state)

For each s_i do

$$T_1[i, 1] = \ln(\pi_i * B_{i,y_1}); T_2[i, 1] = 0;$$

For $i = 2 \dots n$ do

For each s_j do

$$T_1[j, i] = \max_l T_1[l, i - 1] + \ln(A_{l,j} * B_{j,y_i})$$

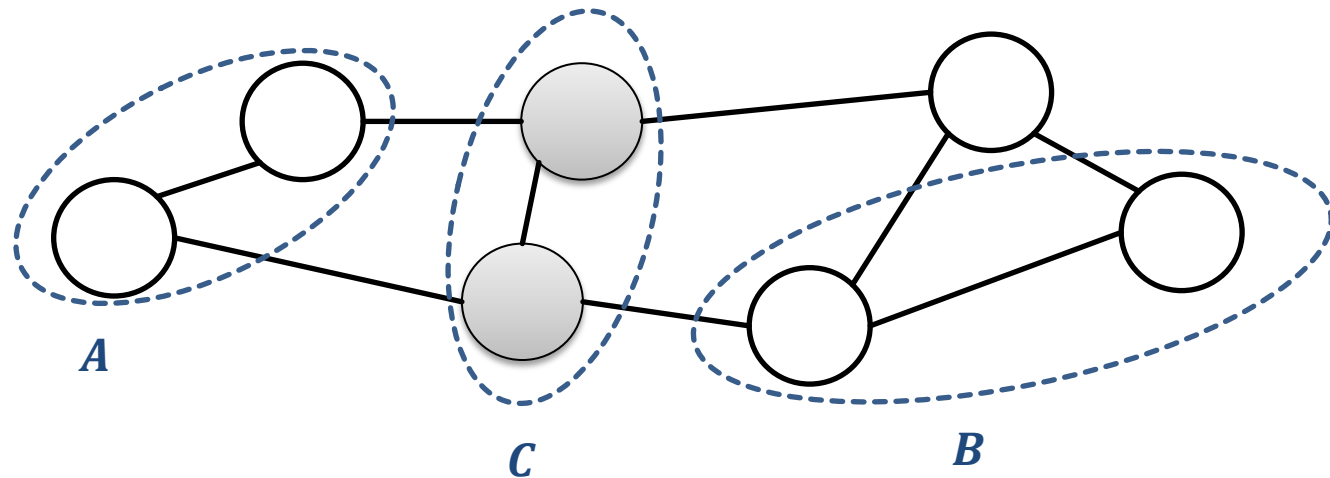
$$T_2[j, i] = \arg \max_l T_1[l, i - 1] + \ln(A_{l,j} * B_{j,y_i})$$

Assign the right state label to each observation by using T_2

Complexity: $O(nk^2)$

Markov random fields

➤ Graph separation



The set of nodes **A** is independent of **B** given **C** if and only if all paths from **A** to **B** lead through **C**; we write: $A \perp B | C$

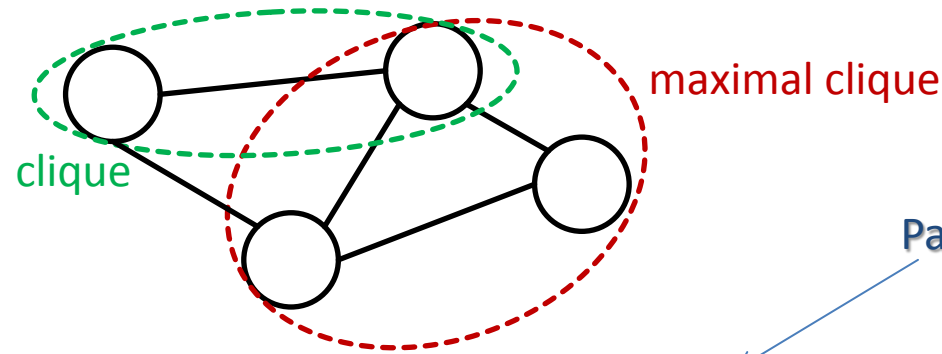
Markov blanket of variable node X is given by all direct neighbors $Nb(X)$ of X

$$P(X_i | X_1, \dots, X_{i-1}, X_{i+1}, \dots, X_n) = P(X_i | Nb(X_i))$$

➔ The joint distribution can be factorized according to above separation rule

Factorization through maximal cliques

- **Observation:** If two nodes X_i, X_j are not directly connected, they are independent given all other nodes → they should **not** occur in same factor



$$P(X_1, \dots, X_n) = \frac{1}{Z} \prod_{C \in \text{MaxCliques}} \psi_C(\mathbf{X}_C)$$

$$Z = \sum_{\mathbf{X}} \prod_{C \in \text{MC}} \psi_C(\mathbf{X}_C)$$

Partition function

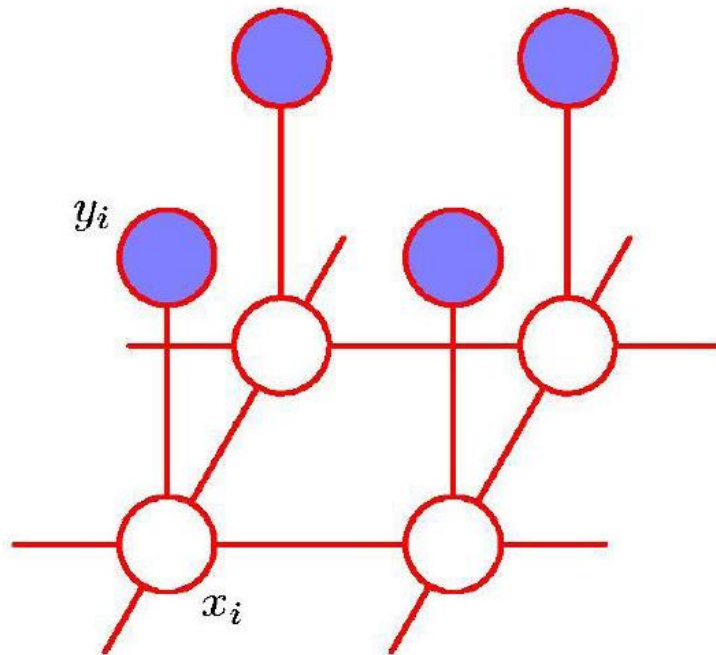
$$\psi_C(\mathbf{X}_C): \text{Potential over clique } C, \text{ typically: } \psi_C(\mathbf{X}_C) = \underbrace{\exp(-E(\mathbf{X}_C))}_{\text{Boltzmann distribution}}$$

Energy function

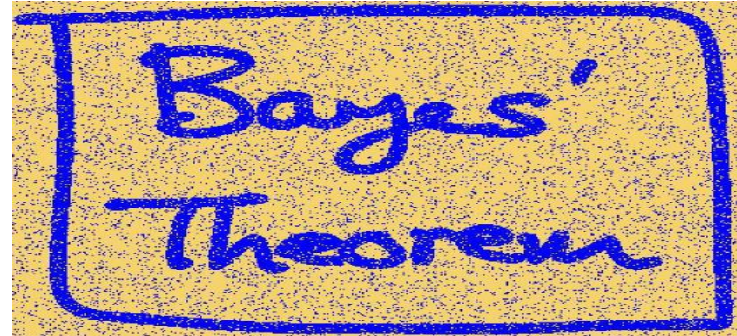
Boltzmann distribution

- **Hammersley-Clifford Theorem:** Factorization through graph separation and factorization through maximal cliques lead to the same sets of distributions

Example: Image de-noising



From C.Bishop: PRML



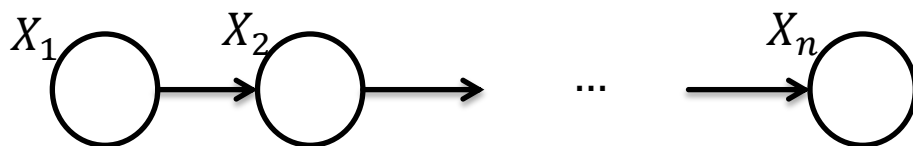
$$E(\mathbf{x}, \mathbf{y}) = h \sum_i x_i - \beta \sum_{\{i,j\}} x_i x_j - \eta \sum_i x_i y_i$$

$$P(\mathbf{x}, \mathbf{y}) = \frac{1}{Z} \exp\{-E(\mathbf{x}, \mathbf{y})\}$$



From directed to undirected chains

- Converting directed into undirected graphical models: chains



$$P(\mathbf{X}) = P(X_1)P(X_2|X_1)P(X_3|X_2) \dots P(X_n|X_{n-1})$$

$$P(\mathbf{X}) = \frac{1}{Z} \psi_{1,2}(X_1, X_2) \psi_{2,3}(X_2, X_3) \dots \psi_{n-1,n}(X_{n-1}, X_n)$$

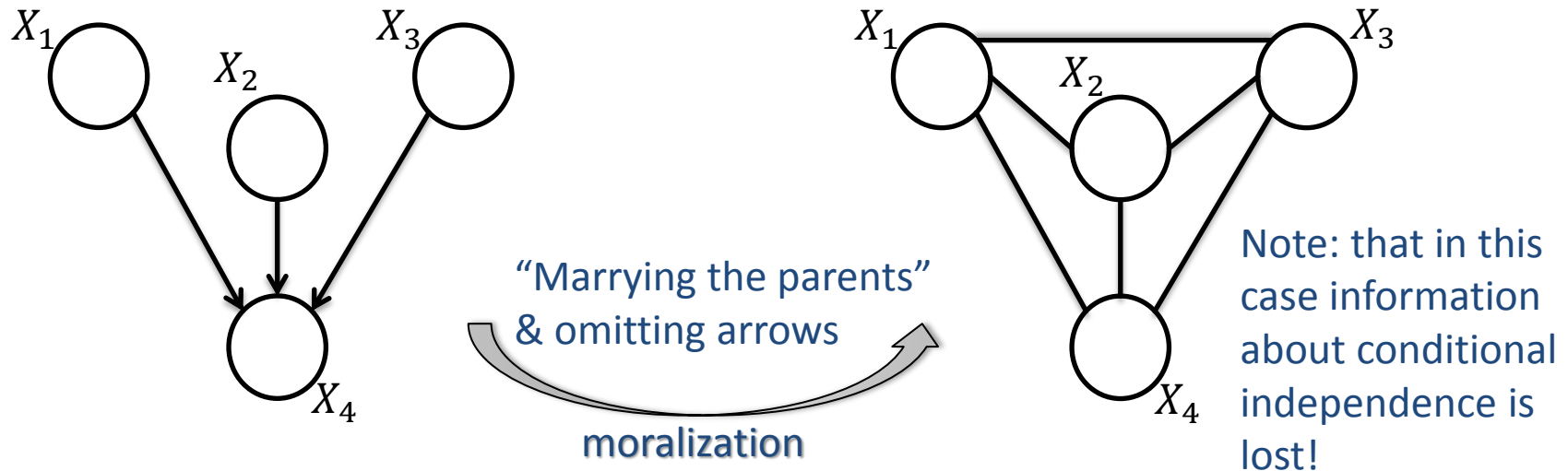


What is the value of the partition function Z ?

Moralization

➤ Converting directed into undirected graphical models: general graphs

Take care that each conditional probability factor in the directed graph is represented by at least one of the maximal cliques in the undirected graph!

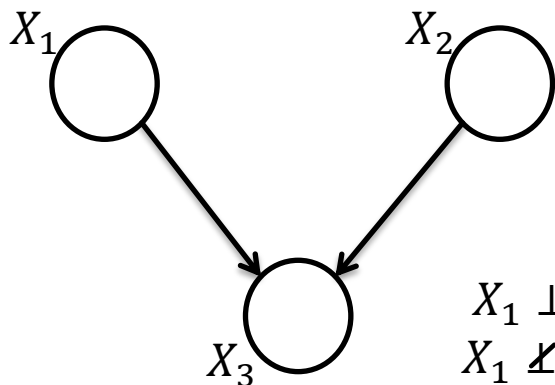
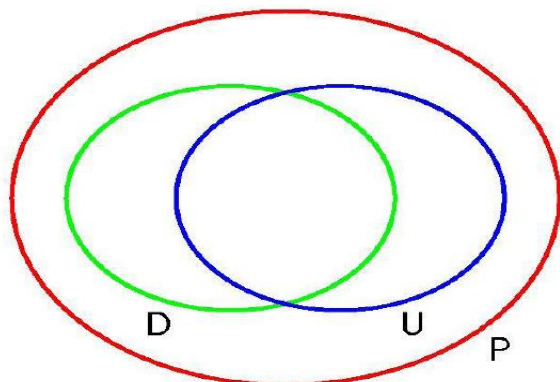


$$P(\mathbf{X}) = P(X_1)P(X_2)P(X_3)P(X_4|X_1, X_2, X_3) = \frac{1}{Z} \psi_{1,2,3,4}(X_1, X_2, X_3, X_4)$$

Theorem: No other technique of turning a directed into an undirected graph retains more independence information than moralization

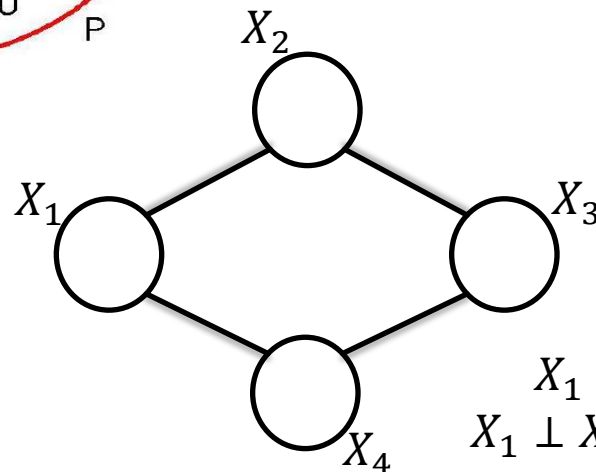
Directed vs. undirected graphical models

From C.Bishop: PRML



$$X_1 \perp X_2 | \emptyset$$

$$X_1 \not\perp X_2 | X_3$$



$$X_1 \not\perp X_2 | \emptyset$$

$$X_1 \perp X_3 | \{X_2, X_3\}$$

$$X_2 \perp X_4 | \{X_1, X_2\}$$

Independence properties cannot be represented by an undirected graph

Independence properties cannot be represented by a directed graph

Summary

- Bayesian networks (directed graphical models)
 - Local Markov property
 - D-separation
 - Distribution filters
 - Example: HMMs

- Markov random fields (undirected graphical models)
 - Graph separation
 - Hammersley-Clifford Theorem
 - Moralization