

Computer Vision Group Prof. Daniel Cremers

Technische Universität München

# 4. Probabilistic Graphical Models Directed Models

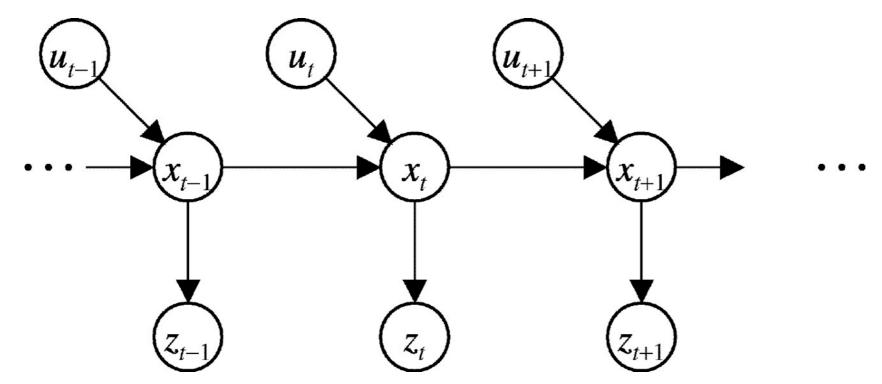
## The Bayes Filter (Rep.)

$$\begin{array}{l} \textbf{Bel}(x_{t}) = p(x_{t} \mid u_{1}, z_{1}, \dots, u_{t}, z_{t}) \\ \textbf{(Bayes)} = \eta \ p(z_{t} \mid x_{t}, u_{1}, z_{1}, \dots, u_{t}) p(x_{t} \mid u_{1}, z_{1}, \dots, u_{t}) \\ \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) p(x_{t} \mid u_{1}, z_{1}, \dots, u_{t}) \\ \textbf{(Tot. prob.)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{1}, z_{1}, \dots, u_{t}, x_{t-1}) \\ p(x_{t-1} \mid u_{1}, z_{1}, \dots, u_{t}) dx_{t-1} \\ \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) p(x_{t-1} \mid u_{1}, z_{1}, \dots, u_{t}) dx_{t-1} \\ \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) p(x_{t-1} \mid u_{1}, z_{1}, \dots, z_{t-1}) dx_{t-1} \\ \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) p(x_{t-1} \mid u_{1}, z_{1}, \dots, z_{t-1}) dx_{t-1} \\ \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(z_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(x_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(x_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(x_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(x_{t} \mid x_{t}) \int p(x_{t} \mid u_{t}, x_{t-1}) \textbf{(Markov)} = \eta \ p(x_{t}$$



## **Graphical Representation (Rep.)**

We can describe the overall process using a Dynamic Bayes Network:



• This incorporates the following Markov assumptions:  $p(z_t \mid x_{0:t}, u_{1:t}, z_{1:t}) = p(z_t \mid x_t)$  (measurement)

$$p(x_t \mid x_{0:t-1}, u_{1:t}, z_{1:t}) = p(x_t \mid x_{t-1}, u_t) \quad \text{(state)}$$



## Definition

A Probabilistic Graphical Model is a diagrammatic representation of a probability distribution.

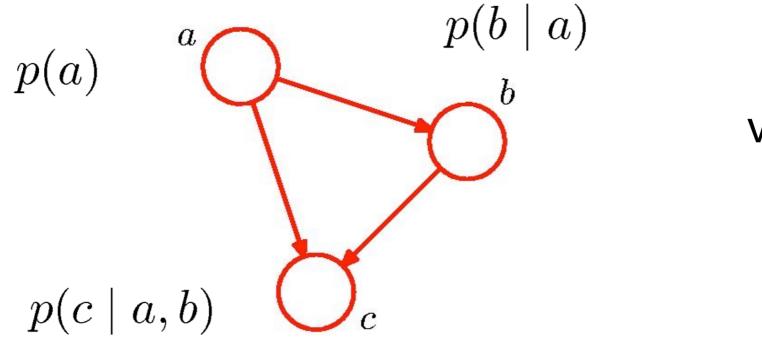
- In a Graphical Model, random variables are represented as nodes, and statistical dependencies are represented using edges between the nodes.
- The resulting graph can have the following properties:
- Cyclic / acyclic
- Directed / undirected
- The simplest graphs are Directed Acyclig Graphs (DAG).





## Simple Example

- Given: 3 random variables a, b, and c
- Joint prob: p(a, b, c) = p(c|a, b)p(a, b) = p(c|a, b)p(b|a)p(a)



Random variables can be discrete or continuous

# A Graphical Model based on a DAG is called a **Bayesian Network**



## Simple Example

In general: K random variables x<sub>1</sub>, x<sub>2</sub>, ..., x<sub>K</sub>
Joint prob:

 $p(x_1, \ldots, x_K) = p(x_K | x_1, \ldots, x_{K-1}) \ldots p(x_2 | x_1) p(x_1)$ 

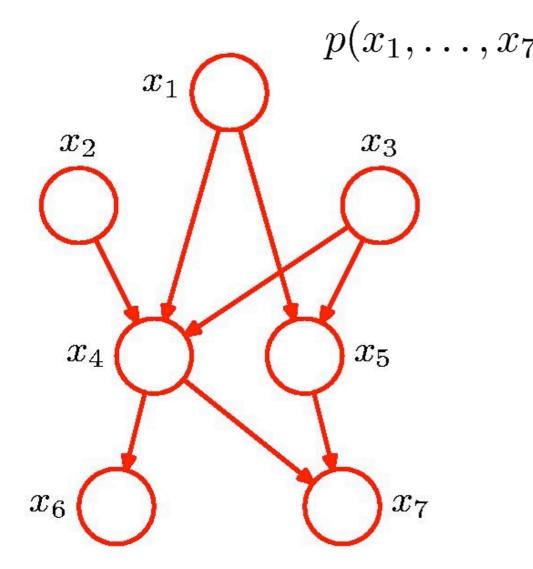
- This leads to a fully connected graph.
- Note: The ordering of the nodes in such a fully connected graph is arbitrary. They all represent the joint probability distribution:

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



## **Bayesian Networks**

Statistical independence can be represented by the absence of edges. This makes the computation efficient.



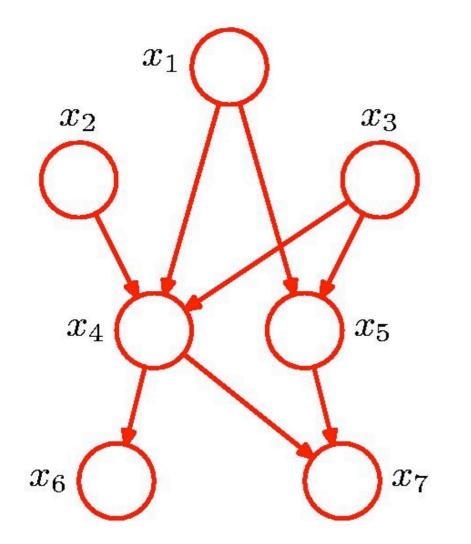
 $p(x_1, \dots, x_7) = p(x_1)p(x_2)p(x_3)p(x_4|x_1, x_2, x_3)$  $p(x_5|x_1, x_3)p(x_6|x_4)p(x_7|x_4, x_5)$ 

Intuitively: only  $_{\mathcal{X}_1}$  and  $_{\mathcal{X}_3}$  have an influence on  $_{\mathcal{X}_5}$ 



## **Bayesian Networks**

We can now define a one-to-one mapping from graphical models to probabilistic formulations:



General Factorization:

$$p(\mathbf{x}) = \prod_{k=1}^{K} p(x_k | \mathrm{pa}_k)$$

where  $pa_k \triangleq ancestors of p_k$ 

and

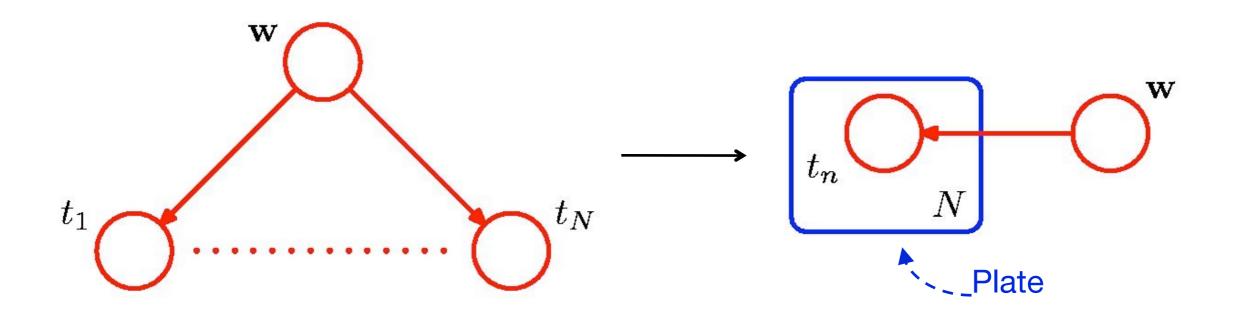
$$p(\mathbf{x}) = p(x_1, \ldots, x_K)$$



## **Elements of Graphical Models**

In case of a series of random variables with equal dependencies, we can subsume them using a **plate:** 

$$p(\mathbf{t}, \mathbf{w}) = p(\mathbf{w}) \prod_{n=1}^{N} p(t_n | \mathbf{w})$$



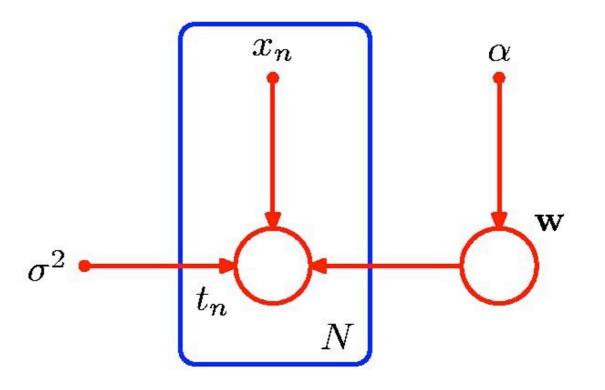


## **Elements of Graphical Models (2)**

We distinguish between **input** variables and explicit **hyper-parameters**:

**A Y** 

$$p(\mathbf{t}, \mathbf{w} | \mathbf{x}, \alpha, \sigma^2) = p(\mathbf{w} | \alpha) \prod_{n=1}^{N} p(t_n | \mathbf{w}, x_n, \sigma^2).$$







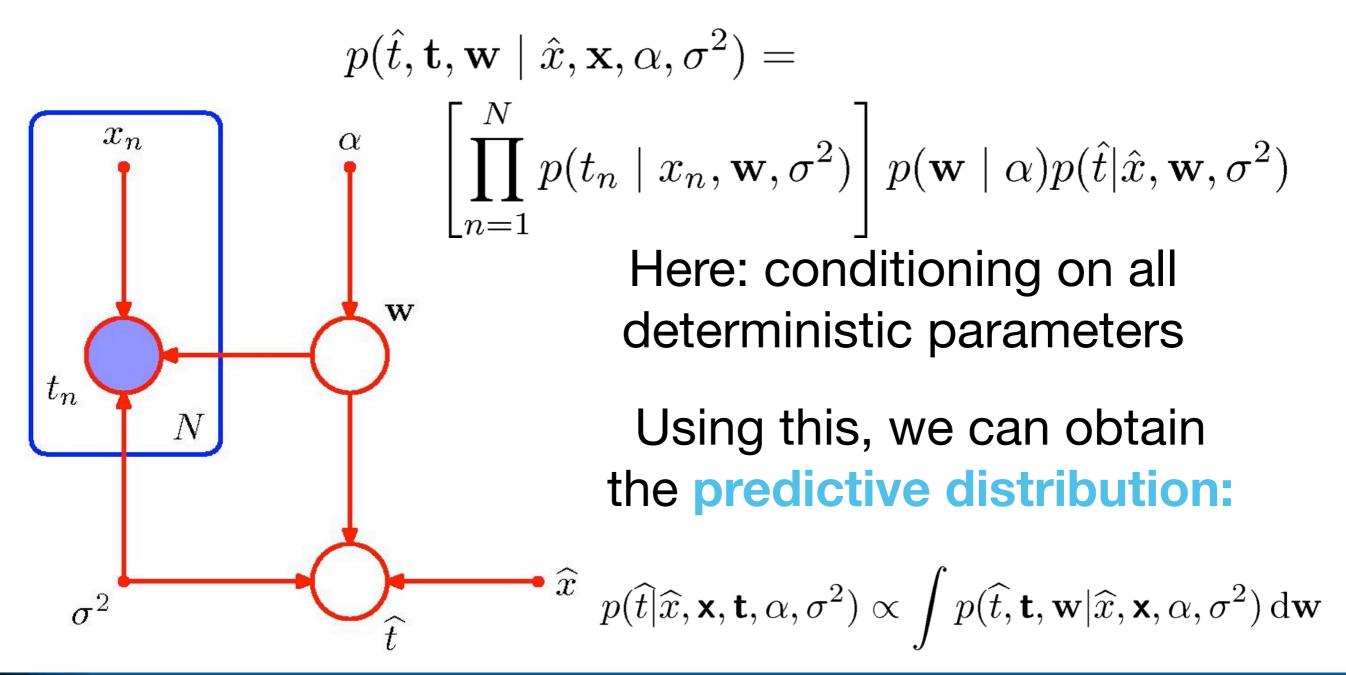
## **Elements of Graphical Models (3)**

We distinguish between **observed** variables and **hidden** variables:



### **Regression as a Graphical Model**

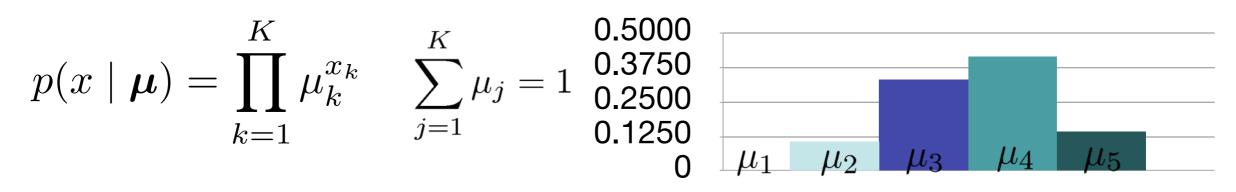
Regression: Prediction of a new target value  $\hat{t}$ 



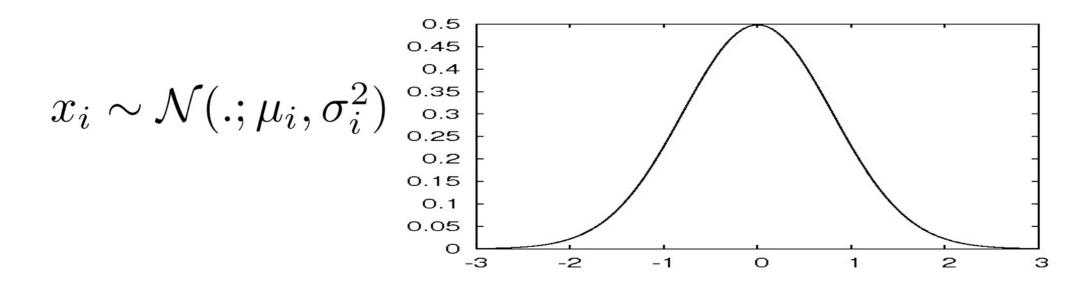


## **Two Special Cases**

- We consider two special cases:
- All random variables are discrete; i.e. Each  $x_i$  is represented by values  $\mu_1, \ldots, \mu_K$  where



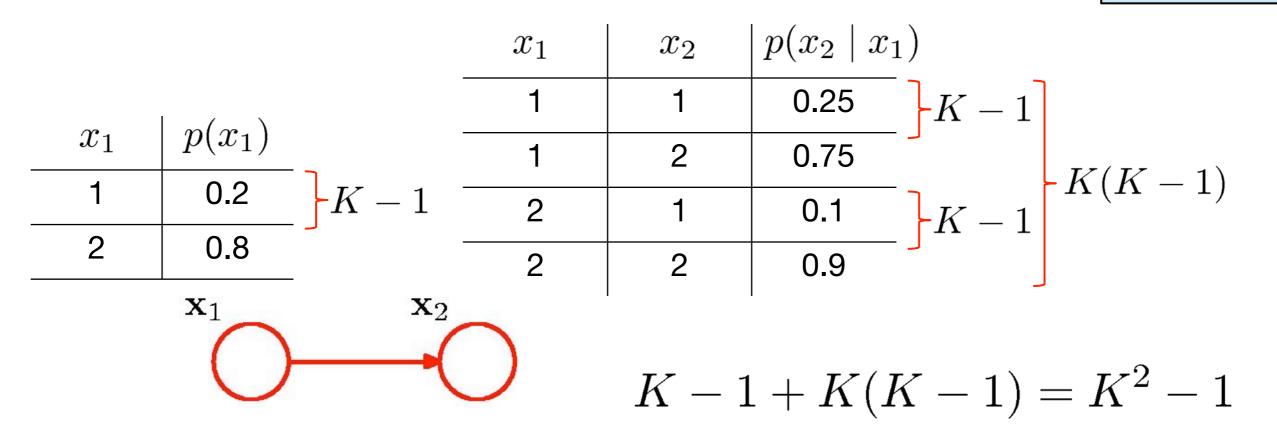
All random variables are Gaussian





## **Discrete Variables: Example**

• Two dependent variables: K<sup>2</sup> - 1 parameters Here: K = 2



Independent joint distribution: 2(K – 1) parameters



$$K - 1 + K - 1 = 2(K - 1)$$



## **Discrete Variables: General Case**

In a general joint distribution with M variables we need to store K<sup>M</sup> -1 parameters

If the distribution can be described by this graph:



then, we have only K -1 + (M -1) K(K -1) parameters.
This graph is called a Markov chain with M nodes.
The number of parameters grows only linearly with the number of variables.



## **Gaussian Variables**

Assume all random variables are Gaussian and we define

$$p(x_i \mid \mathrm{pa}_i) = \mathcal{N}\left(x_i; \sum_{j \in \mathrm{pa}_i} w_{ij} x_j + b_i, v_i\right)$$

Then one can show that the joint probability p(x) is a multivariate Gaussian. Furthermore:

$$x_i = \sum_{j \in pa_i} w_{ij} x_j + b_j + \sqrt{v_i} \epsilon_i \qquad \epsilon_i \sim \mathcal{N}(0, 1)$$

Thus:

$$E[x_i] = \sum_{j \in pa_i} w_{ij} E[x_j] + b_i$$

i.e., we can compute the mean values recursively.



## **Gaussian Variables**

Assume all random variables are Gaussian and we define

$$p(x_i \mid \mathrm{pa}_i) = \mathcal{N}\left(x_i; \sum_{i \in \mathrm{pa}_i} w_{ij}x_j + b_i, v_i\right)$$

The same can be shown for the covariance. Thus:

Mean and covariance can be calculated recursively

Furthermore it can be shown that:

- The fully connected graph corresponds to a Gaussian with a general symmetric covariance matrix
- The non-connected graph corresponds to a diagonal covariance matrix



## Independence (Rep.)

**Definition 1.4:** Two random variables X and Y are *independent* iff: p(x, y) = p(x)p(y)

For independent random variables  $\chi$  and  $\gamma$  we have:

$$p(x \mid y) = \frac{p(x, y)}{p(y)} = \frac{p(x)p(y)}{p(y)} = p(x)$$

Notation:	$x \perp\!\!\!\perp y \mid \emptyset$
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Independence does not imply conditional independence. The same is true for the opposite case.



## **Conditional Independence (Rep.)**

**Definition 1.5:** Two random variables X and Y are conditional independent given a third random variable Z iff:

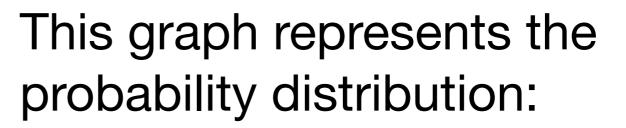
$$p(x, y \mid z) = p(x \mid z)p(y \mid z)$$

This is equivalent to:

$$p(x \mid z) = p(x \mid y, z) \text{ and}$$
$$p(y \mid z) = p(y \mid x, z)$$

Notation: 
$$x \perp \!\!\!\perp y \mid z$$





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

Marginalizing out *c* on both sides gives

$$p(a,b) = \sum_{c} p(a|c)p(b|c)p(c)$$

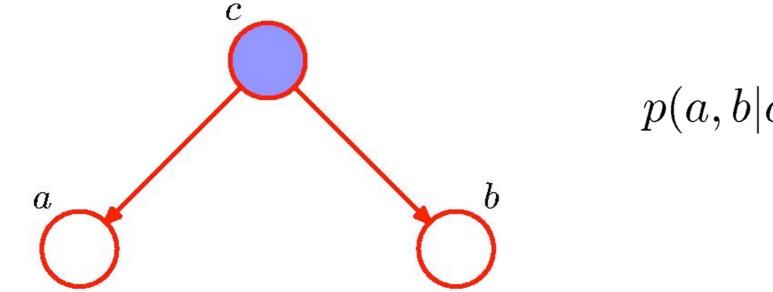
This is in general not equal to p(a)p(b).

**Thus:** *a* and *b* are not independent:  $a \not\!\!\!\perp b \mid \emptyset$ 

a



• Now, we condition on c (it is assumed to be known):

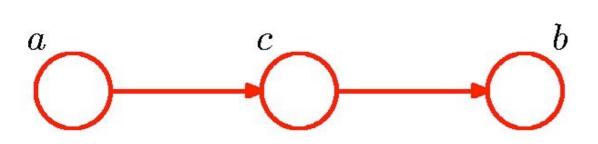


$$(a,b|c) = \frac{p(a,b,c)}{p(c)}$$
$$= p(a|c)p(b|c)$$

**Thus:** *a* and *b* are conditionally independent given *c*:  $a \perp b \mid c$ We say that the node at *c* is a **tail-to-tail node** on the path between *a* and *b* 







This graph represents the distribution:

p(a, b, c) = p(a)p(c|a)p(b|c)

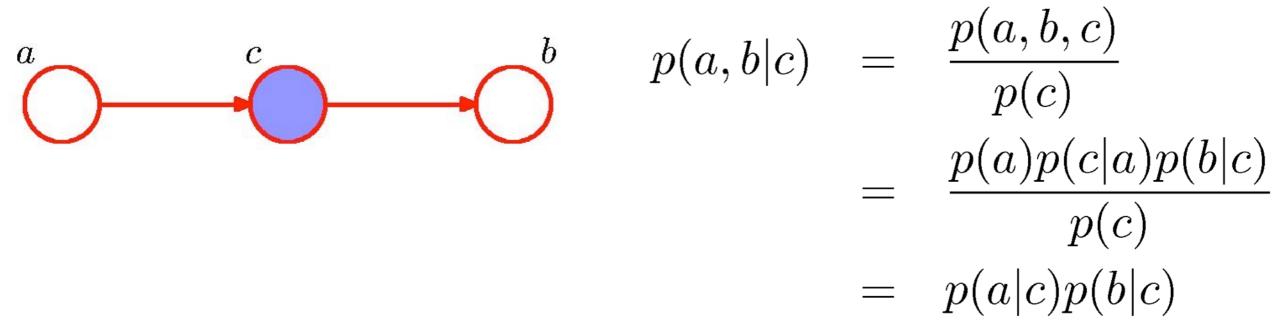
Again, we marginalize over c:

$$p(a,b) = p(a) \sum_{c} p(c|a)p(b|c) = p(a) \sum_{c} p(c|a)p(b|c,a)$$
$$= p(a) \sum_{c} \frac{p(c,a)p(b,c,a)}{p(a)p(c,a)} = p(a) \sum_{c} p(b,c \mid a)$$
$$= p(a)p(b|a)$$

And we obtain:  $a \not\perp b \mid \emptyset$ 



As before, now we condition on c:



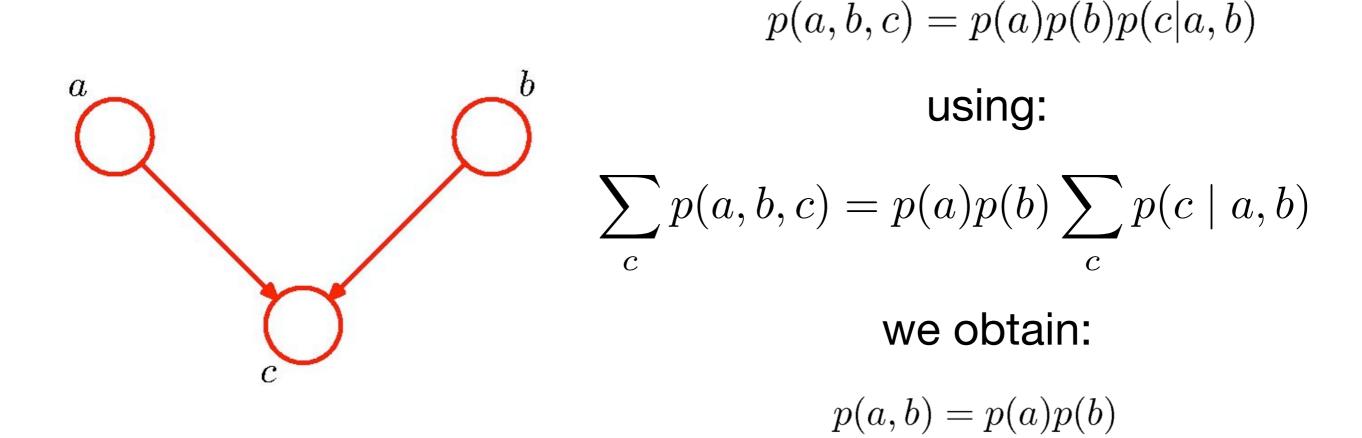
And we obtain:  $a \perp b \mid c$ 

#### We say that the node at c is a head-to-tail node on the path between a and b.





Now consider this graph:

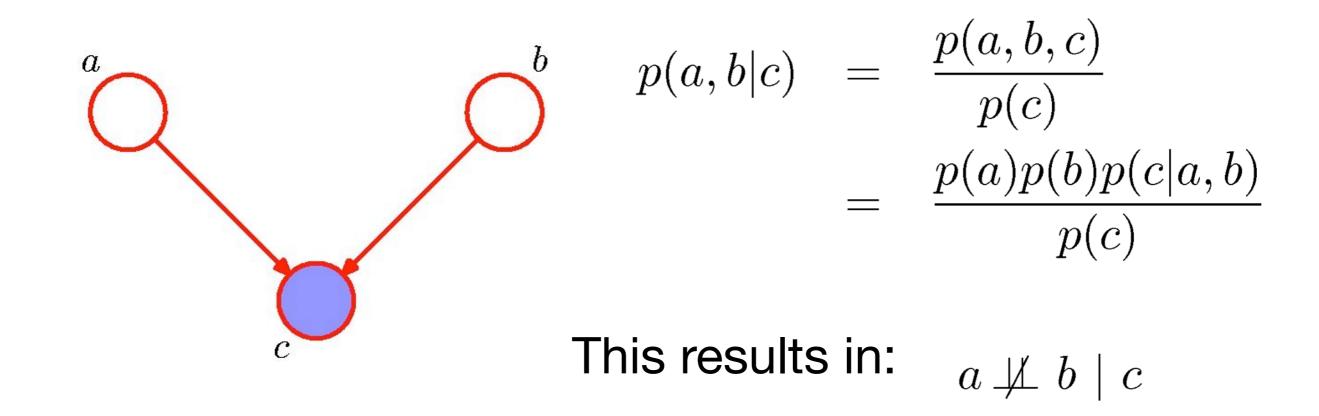


#### And the result is: $a \perp b \mid \emptyset$





#### Again, we condition $on_c$



We say that the node at c is a head-to-head node on the path between a and b.



## **To Summarize**

- When does the graph represent (conditional) independence?
  - Tail-to-tail case: if we condition on the tail-to-tail node Head-to-tail case: if we cond. on the head-to-tail node Head-to-head case: if we do not condition on the headto-head node (and neither on any of its descendants)

In general, this leads to the notion of D-separation for directed graphical models.





## **D-Separation**

Say: A, B, and C are non-intersecting subsets of nodes in a directed graph.

- A path from A to B is blocked by C if it contains a node such that either
- a) the arrows on the path meet either head-to-tail or tail-totail at the node, and the node is in the set C, or
- b) the arrows meet **head-to-head** at the node, and neither the node, nor any of its descendants, are in the set C.
- If all paths from A to B are blocked, A is said to be d-separated from B by C.
- Notation: dsep(A, B|C)

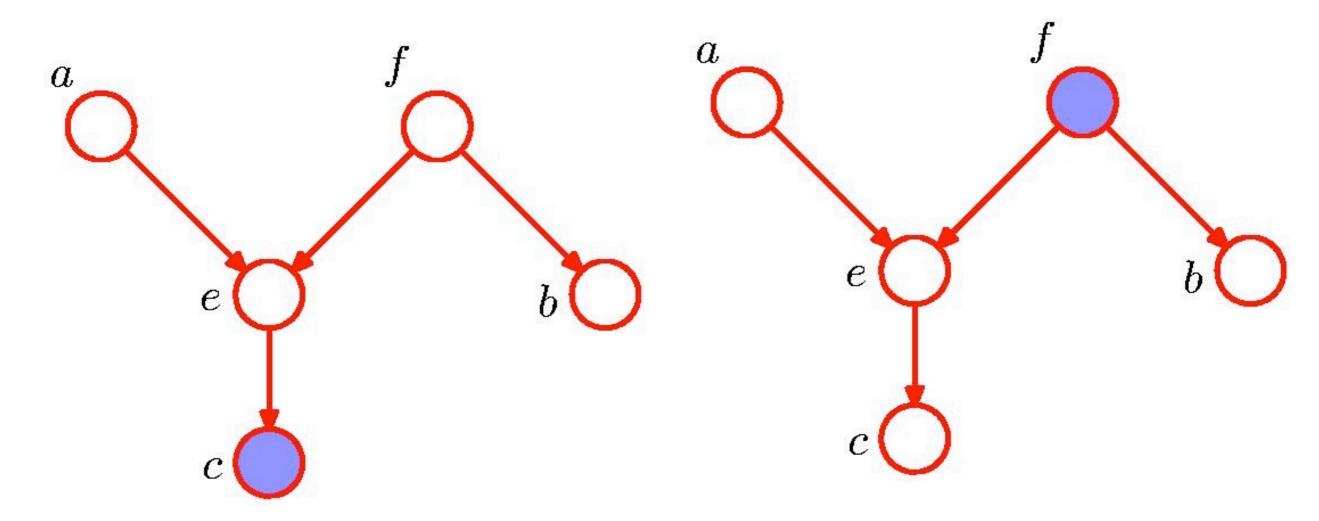


## **D-Separation**

Say: A, B, and C are non-intersecting subsets of **D-Separation is a** nodes A path ntains property of graphs a nod a) the a <sup>r</sup> tail-toand not of tail at t probability b) the a neither the noc **J**. distributions If all p aid to be d-separated from B by C. Notation: dsep(A, B|C)



## **D-Separation: Example**



#### $\neg \operatorname{dsep}(a, b|c)$

We condition on a descendant of e, i.e. it does not block the path from a to b.

#### $\operatorname{dsep}(a, b|f)$

We condition on a tail-to-tail node on the only path from a to b, i.e f blocks the path.





## I-Map

**Definition 4.1:** A graph G is called an I-map for a distribution p if every D-separation of G corresponds to a conditional independence relation satisfied by p:

### $\forall A, B, C : \operatorname{dsep}(A, B, C) \Rightarrow A \perp\!\!\!\perp B \mid C$

**Example:** The fully connected graph is an I-map for any distribution, as there are no D-separations in that graph.





## **D-Map**

**Definition 4.2:** A graph G is called an **D-map** for a distribution p if for every conditional independence relation satisfied by p there is a D-separation in G :

## $\forall A, B, C : A \perp\!\!\!\perp B \mid C \Rightarrow \operatorname{dsep}(A, B, C)$

**Example:** The graph without any edges is a D-map for any distribution, as all pairs of subsets of nodes are D-separated in that graph.





## **Perfect Map**

**Definition 4.3:** A graph G is called a perfect map for a distribution p if it is a D-map and an I-map of p.

## $\forall A, B, C : A \perp\!\!\!\perp B \mid C \Leftrightarrow \operatorname{dsep}(A, B, C)$

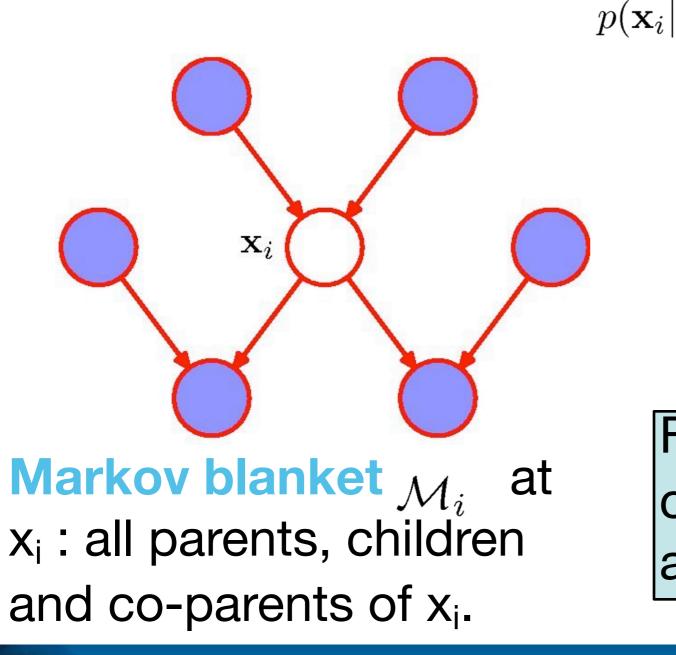
A perfect map uniquely defines a probability distribution.





## **The Markov Blanket**

 Consider a distribution of a node x\_i conditioned on all other nodes:



$$\mathbf{x}_{\{j \neq i\}}) = \frac{p(\mathbf{x}_1, \dots, \mathbf{x}_M)}{\int p(\mathbf{x}_1, \dots, \mathbf{x}_M) d\mathbf{x}_i}$$
$$= \frac{\prod_k p(\mathbf{x}_k | \mathbf{pa}_k)}{\int \prod_k p(\mathbf{x}_k | \mathbf{pa}_k) d\mathbf{x}_i}$$
$$= p(\mathbf{x}_i | \mathbf{x}_{\mathcal{M}_i})$$

Factors independent of x<sub>i</sub> cancel between numerator and denominator.





## Summary

- Graphical models represent joint probability distributions using nodes for the random variables and edges to express (conditional) (in)dependence
- A prob. dist. can always be represented using a fully connected graph, but this is inefficient
- In a directed acyclic graph, conditional independence is determined using D-separation
- A perfect map implies a one-to-one mapping between c.i. relations and D-separations
- The Markov blanket is the minimal set of observed nodes to obtain conditional independence



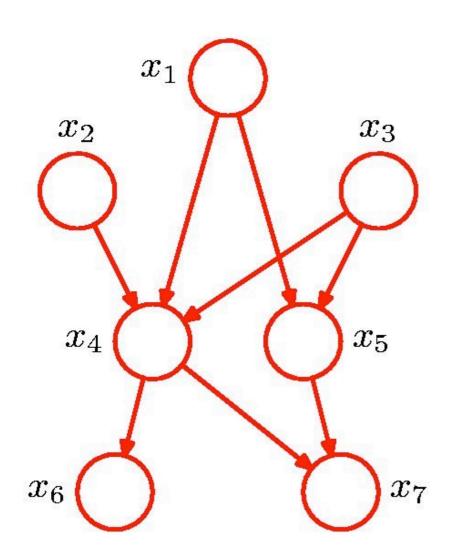


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# 4. Probabilistic Graphical Models Undirected Models

## **Repetition: Bayesian Networks**

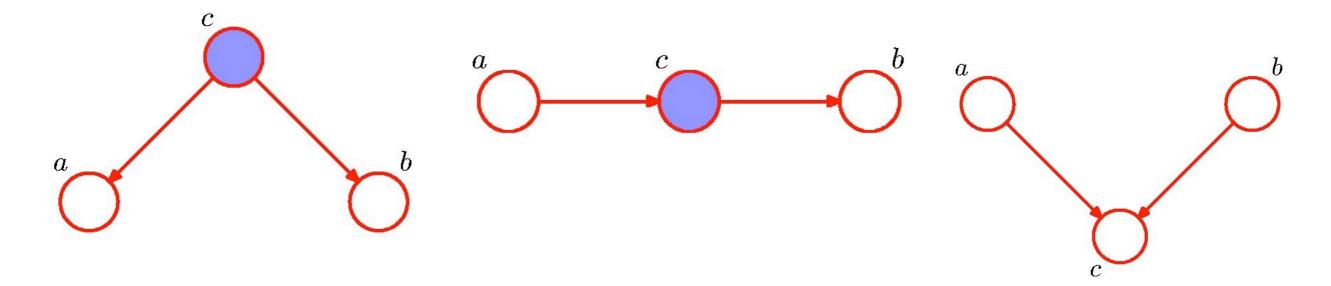


Directed graphical models can be used to represent **probability distributions** This is useful to do **inference** and to **generate samples** from the distribution efficiently

$$p(x_1, \dots, x_7) = p(x_1)p(x_2)p(x_3)p(x_4|x_1, x_2, x_3)$$
$$p(x_5|x_1, x_3)p(x_6|x_4)p(x_7|x_4, x_5)$$



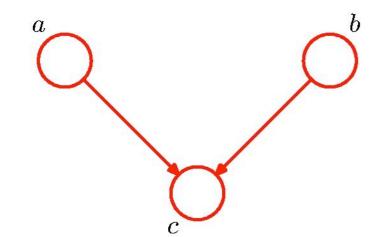
# **Repetition: D-Separation**



- D-separation is a property of graphs that can be easily determined
- An I-map assigns every d-separation a c.i. rel
- A D-map assigns every c.i. rel a d-separation
- Every Bayes net determines a unique prob. dist.



# In-depth: The Head-to-Head Node



$$p(a) = 0.9 \qquad p(b) = 0.9$$

$$a \qquad b \qquad p(c)$$

$$1 \qquad 1 \qquad 0.8$$

$$1 \qquad 0 \qquad 0.2$$

$$0 \qquad 1 \qquad 0.2$$

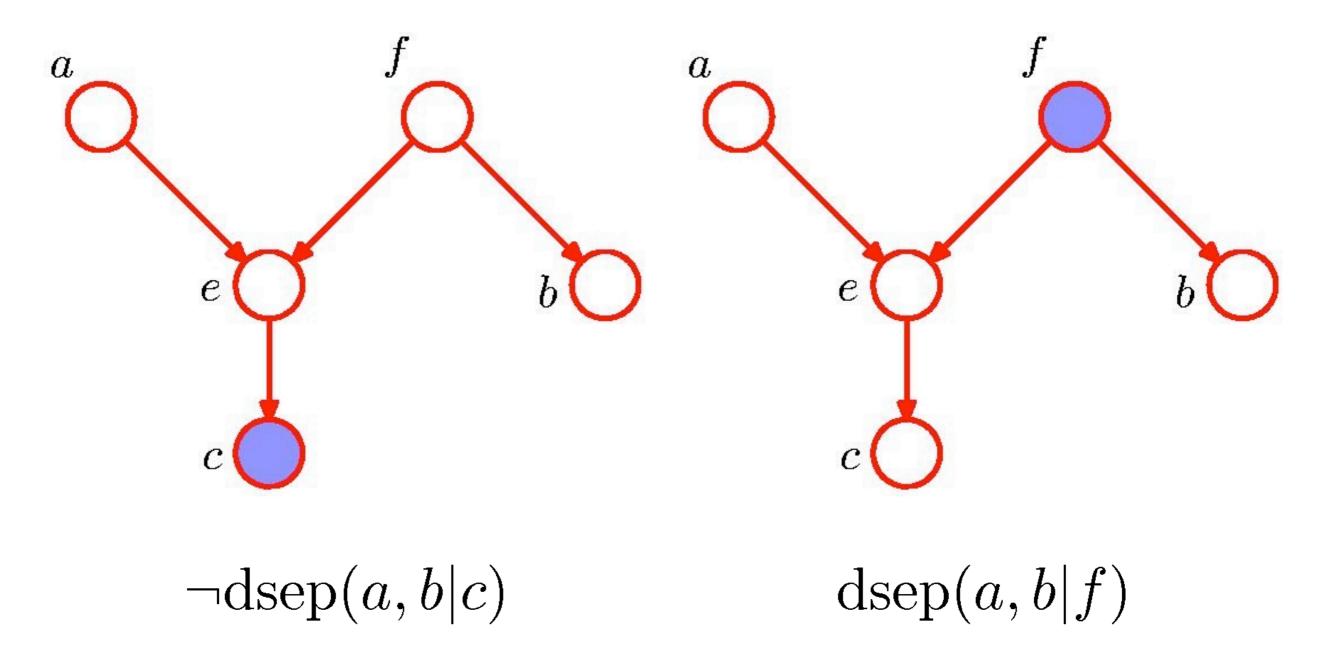
$$0 \qquad 0 \qquad 1$$

Example:

- a: Battery charged (0 or 1)
- b: Fuel tank full (0 or 1)
- c: Fuel gauge says full (0 or 1)
- We can compute  $p(\neg c) = 0.315$
- **and**  $p(\neg c \mid \neg b) = 0.81$
- and obtain  $p(\neg b \mid \neg c) \approx 0.257$
- similarly:  $p(\neg b \mid \neg c, \neg a) \approx 0.111$
- "*a* explains *c* away"



# **Repetition: D-Separation**





## **Directed vs. Undirected Graphs**

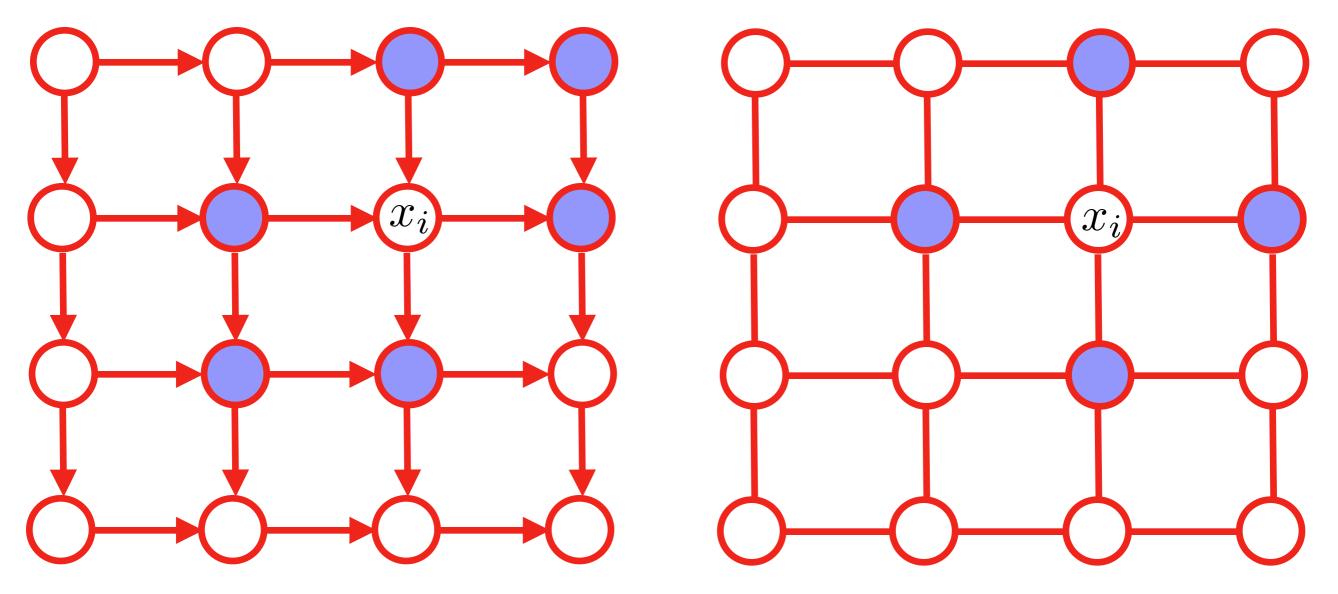
Using D-separation we can identify conditional independencies in directed graphical models, but:

- Is there a simpler, more intuitive way to express conditional independence in a graph?
- Can we find a representation for cases where an "ordering" of the random variables is inappropriate (e.g. the pixels in a camera image)?

Yes, we can: by removing the directions of the edges we obtain an Undirected Graphical Model, also known as a Markov Random Field



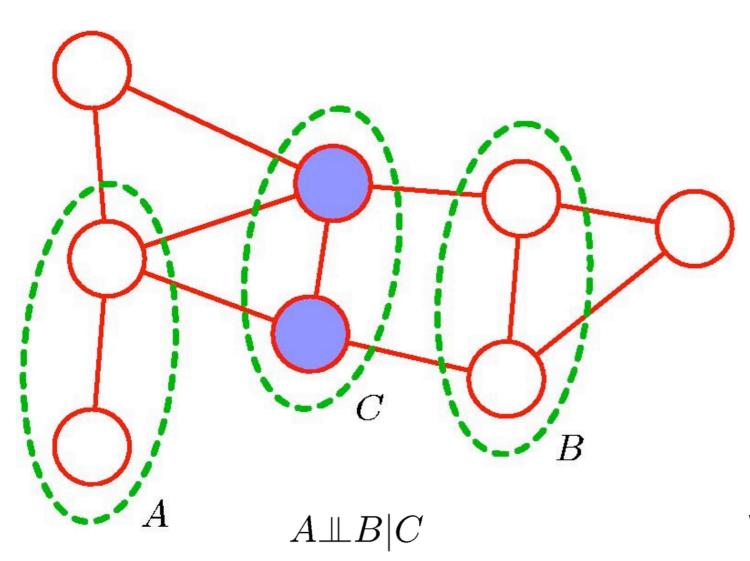
# **Example: Camera Image**



- directions are counter-intuitive for images
- Markov blanket is not just the direct neighbors when using a directed model



### **Markov Random Fields**



All paths from *A* to *B* go through *C*, i.e. *C* blocks all paths.

Markov Blanket

We only need to condition on the **direct neighbors** of

x to get c.i., because these already block every path from x to any other node.



# **Factorization of MRFs**

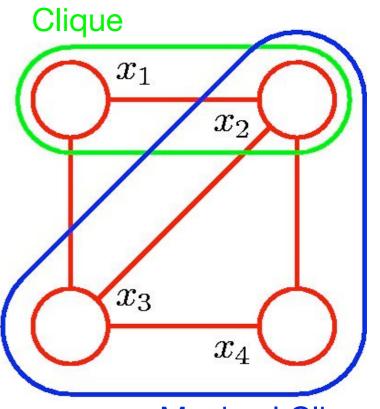
Any two nodes  $x_i$  and  $x_j$  that are not connected in an MRF are conditionally independent given all other nodes:

 $p(x_i, x_j \mid \mathbf{x}_{\backslash \{i, j\}}) = p(x_i \mid \mathbf{x}_{\backslash \{i, j\}}) p(x_j \mid \mathbf{x}_{\backslash \{i, j\}})$ 

In turn: each factor contains only nodes that are connected

This motivates the consideration of cliques in the graph:

- A clique is a fully connected subgraph.
- A maximal clique can not be extended with another node without loosing the property of full connectivity.



#### **Maximal Clique**



## **Factorization of MRFs**

In general, a Markov Random Field is factorized as

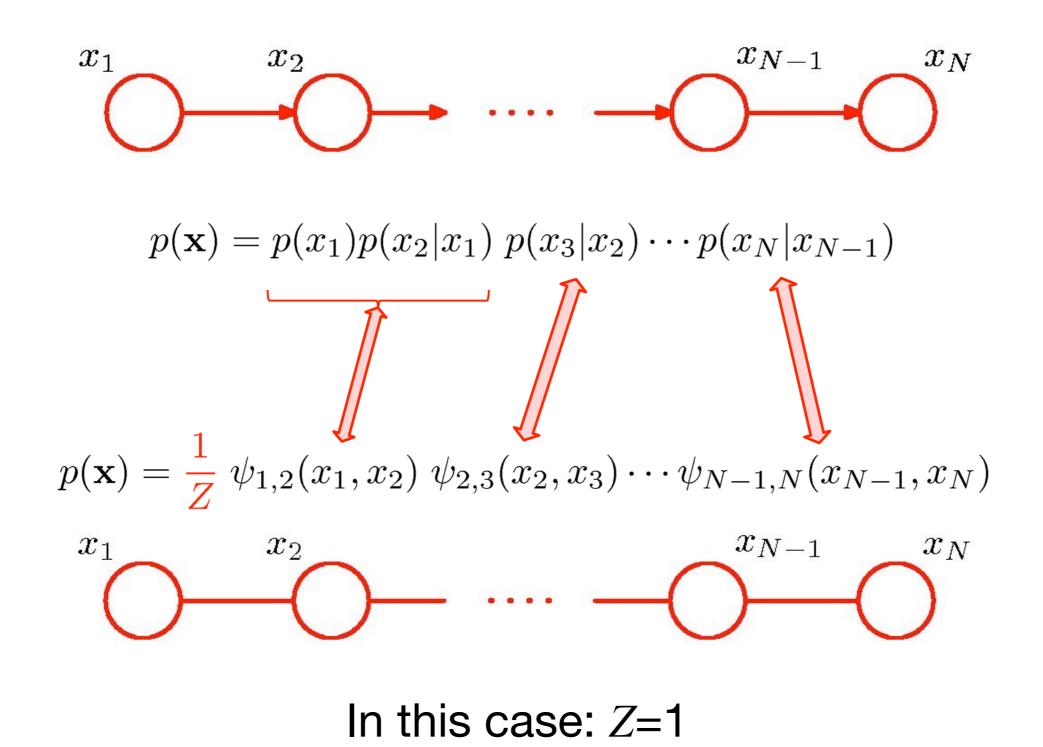
$$p(\mathbf{x}) = \frac{\prod_C \phi_C(\mathbf{x}_C)}{\sum_{\mathbf{x}'} \prod_C \phi_C(\mathbf{x}'_C)} = \frac{1}{Z} \prod_C \phi_C(\mathbf{x}_C)$$
(4.1)

where *C* is the set of all (maximal) cliques and  $\Phi_C$  is a positive function of a given clique  $\mathbf{x}_C$  of nodes, called the **clique potential**. *Z* is called the **partition function**. **Theorem (Hammersley/Clifford):** Any undirected model with associated clique potentials  $\Phi_C$  is a perfect map for the probability distribution defined by Equation (4.1).

As a conclusion, all probability distributions that can be factorized as in (4.1), can be represented as an MRF.

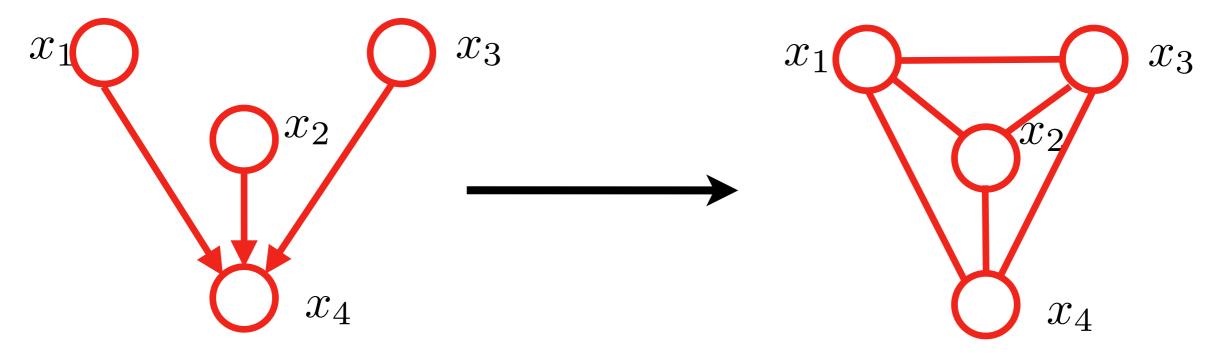


#### **Converting Directed to Undirected Graphs (1)**





## **Converting Directed to Undirected Graphs (2)**



$$p(\mathbf{x}) = p(x_1)p(x_2)p(x_2)p(x_4 \mid x_1, x_2, x_3)$$

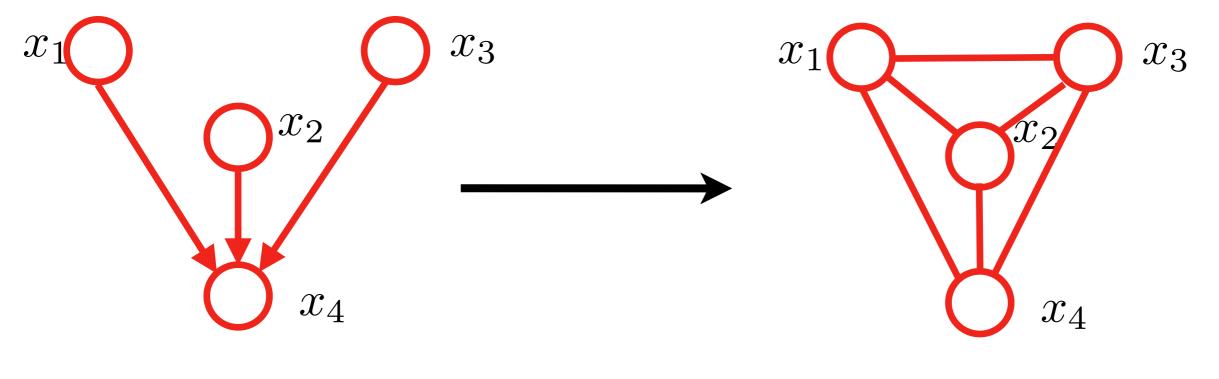
In general: conditional distributions in the directed graph are mapped to cliques in the undirected graph

**However:** the variables are **not** conditionally independent given the head-to-head node

Therefore: Connect all parents of head-to-head nodes with each other (moralization)



#### **Converting Directed to Undirected Graphs (2)**



 $p(\mathbf{x}) = p(x_1)p(x_2)p(x_2)p(x_4 \mid x_1, x_2, x_3)$ 

 $p(\mathbf{x}) = \phi(x_1, x_2, x_3, x_4)$ 

**Problem:** This process can remove conditional independence relations (inefficient)

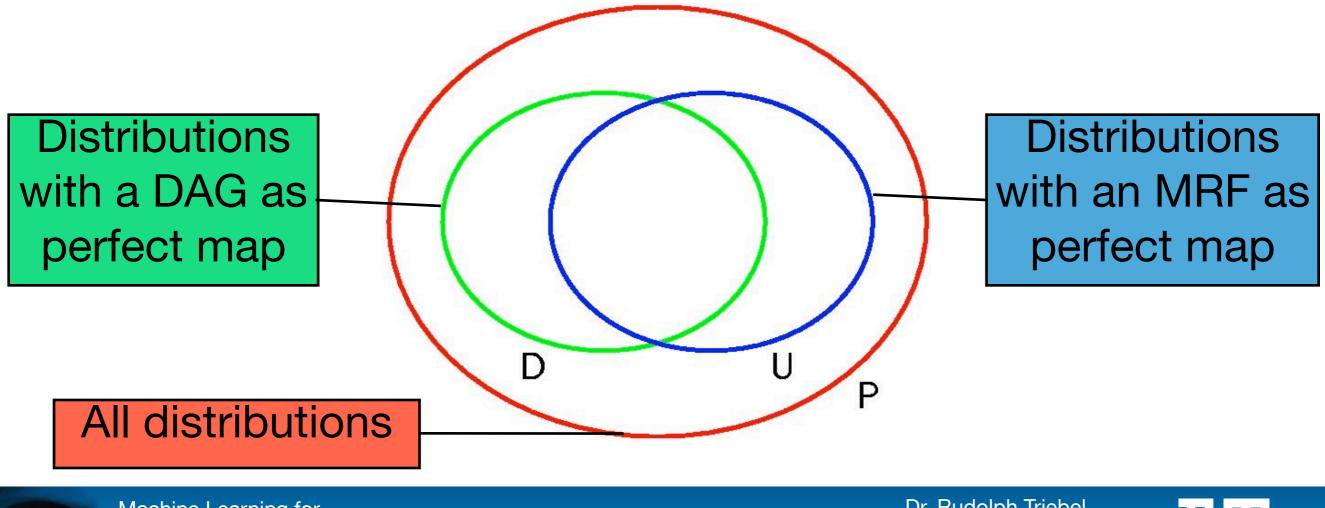
**Generally:** There is no one-to-one mapping between the distributions represented by directed and by undirected graphs.





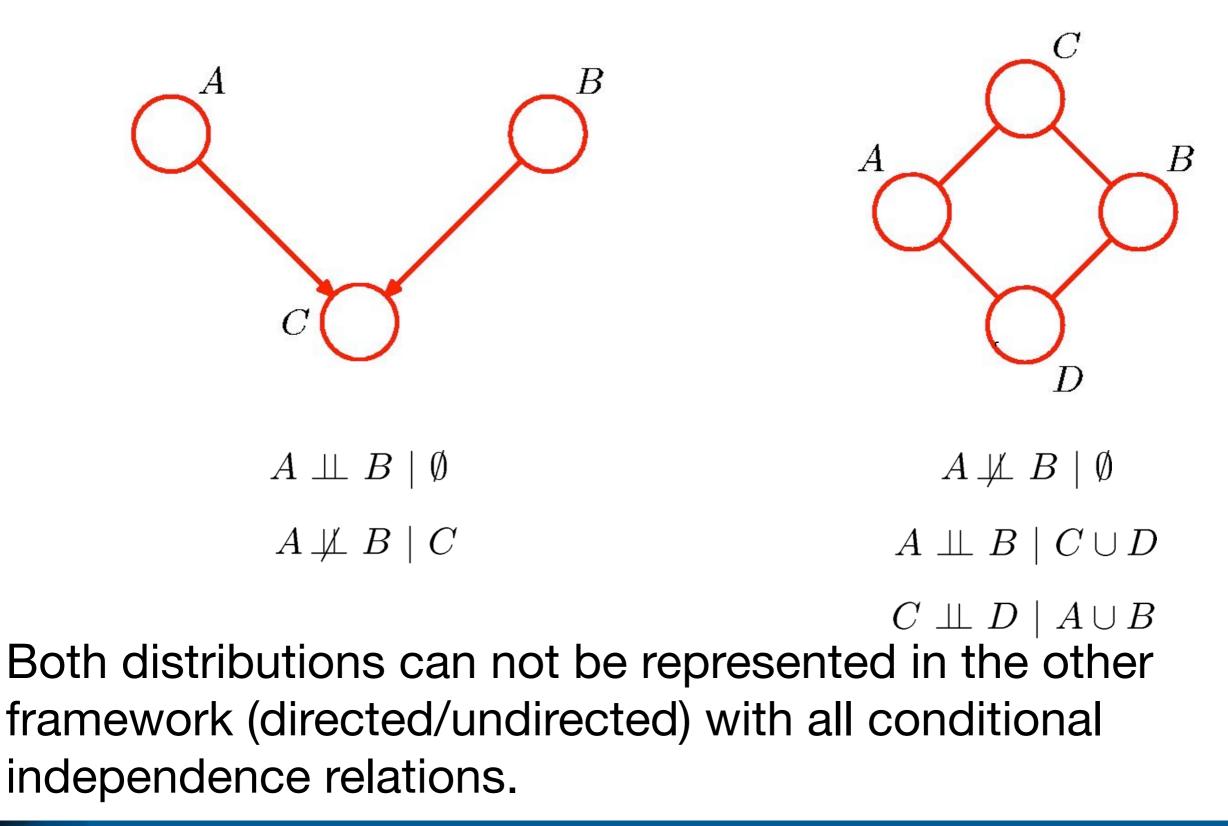
## Representability

- As for DAGs, we can define an I-map, a D-map and a perfect map for MRFs.
- The set of all distributions for which a DAG exists that is a perfect map is different from that for MRFs.





#### **Directed vs. Undirected Graphs**





# **Using Graphical Models**

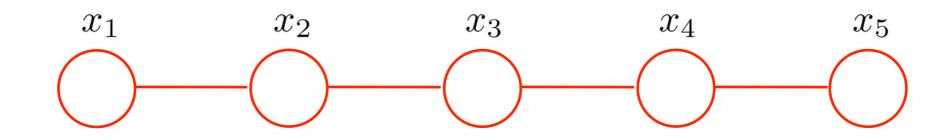
We can use a graphical model to do inference:

- Some nodes in the graph are observed, for others we want to find the posterior distribution
- Also, computing the local marginal distribution p(x<sub>n</sub>) at any node x<sub>n</sub> can be done using inference.

Question: How can inference be done with a graphical model?

We will see that when exploiting conditional independences we can do efficient inference.





The joint probability is given by

$$p(\mathbf{x}) = \frac{1}{Z} \psi_{1,2}(x_1, x_2) \psi_{2,3}(x_2, x_3) \psi_{3,4}(x_3, x_4) \psi_{4,5}(x_4, x_5)$$

The marginal at  $x_3$  is  $p(x_3) = \sum_{x_1} \sum_{x_2} \sum_{x_4} \sum_{x_5} p(\mathbf{x})$ 

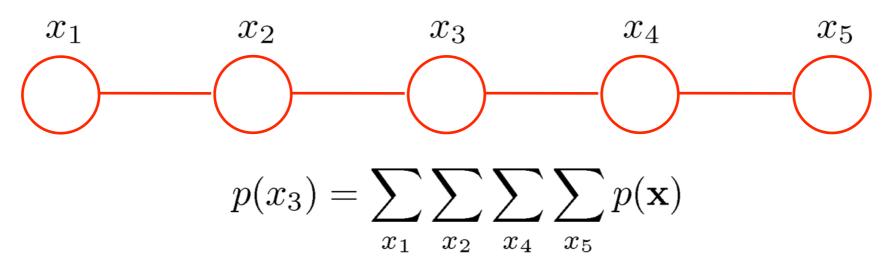
In the general case with N nodes we have

$$p(\mathbf{x}) = \frac{1}{Z} \psi_{1,2}(x_1, x_2) \psi_{2,3}(x_2, x_3) \cdots \psi_{N-1,N}(x_{N-1}, x_N)$$

and

$$p(x_n) = \sum_{x_1} \cdots \sum_{x_{n-1}} \sum_{x_{n+1}} \cdots \sum_{x_N} p(\mathbf{x})$$





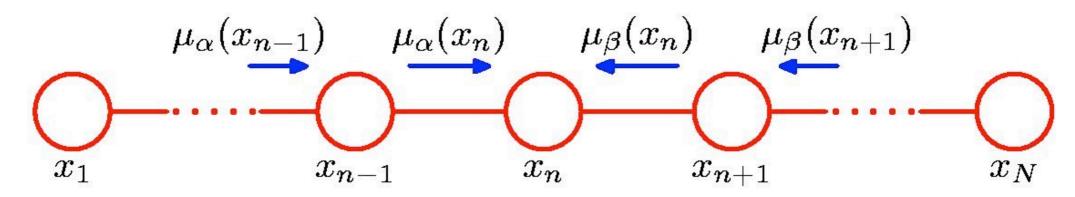
 This would mean K<sup>N</sup> computations! A more efficient way is obtained by rearranging:

$$p(x_{3}) = \frac{1}{Z} \sum_{x_{1}} \sum_{x_{2}} \sum_{x_{4}} \sum_{x_{5}} \psi_{1,2}(x_{1}, x_{2})\psi_{2,3}(x_{2}, x_{3})\psi_{3,4}(x_{3}, x_{4})\psi_{4,5}(x_{4}, x_{5})$$

$$= \frac{1}{Z} \sum_{x_{2}} \sum_{x_{1}} \sum_{x_{4}} \sum_{x_{5}} \psi_{1,2}(x_{1}, x_{2})\psi_{2,3}(x_{2}, x_{3})\psi_{3,4}(x_{3}, x_{4})\psi_{4,5}(x_{4}, x_{5})$$

$$= \frac{1}{Z} \sum_{x_{2}} \psi_{2,3}(x_{2}, x_{3}) \sum_{x_{1}} \psi_{1,2}(x_{1}, x_{2}) \sum_{x_{4}} \psi_{3,4}(x_{3}, x_{4}) \sum_{x_{5}} \psi_{4,5}(x_{4}, x_{5})$$

$$\mu_{\alpha}(x_{3}) \leftarrow \text{Vectors of size K} \rightarrow \mu_{\beta}(x_{3})$$



In general, we have

$$p(x_n) = \frac{1}{Z} \left[ \sum_{x_{n-1}} \psi_{n-1,n}(x_{n-1}, x_n) \cdots \left[ \sum_{x_1} \psi_{1,2}(x_1, x_2) \right] \cdots \right] \\ \mu_{\alpha}(x_n) \\ \left[ \sum_{x_{n+1}} \psi_{n,n+1}(x_n, x_{n+1}) \cdots \left[ \sum_{x_N} \psi_{N-1,N}(x_{N-1}, x_N) \right] \cdots \right] \\ \mu_{\beta}(x_n)$$

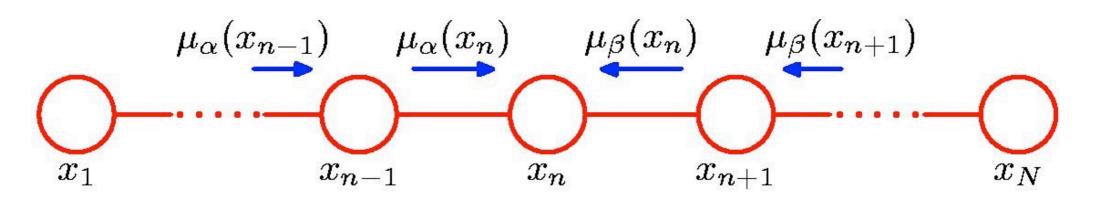


The messages  $\mu_{\alpha}$  and  $\mu_{\beta}$  can be computed recursively:

$$\mu_{\alpha}(x_{n}) = \sum_{x_{n-1}} \psi_{n-1,n}(x_{n-1}, x_{n}) \left[ \sum_{x_{n-2}} \cdots \right]$$
$$= \sum_{x_{n-1}} \psi_{n-1,n}(x_{n-1}, x_{n}) \mu_{\alpha}(x_{n-1}).$$
$$\mu_{\beta}(x_{n}) = \sum_{x_{n+1}} \psi_{n,n+1}(x_{n}, x_{n+1}) \left[ \sum_{x_{n+2}} \cdots \right]$$
$$= \sum_{x_{n+1}} \psi_{n,n+1}(x_{n}, x_{n+1}) \mu_{\beta}(x_{n+1}).$$

Computation of  $\mu_{\alpha}$  starts at the first node and computation of  $\mu_{\beta}$  starts at the last node.





• The first values of  $\mu_{\alpha}$  and  $\mu_{\beta}$  are:

$$\mu_{\alpha}(x_2) = \sum_{x_1} \psi_{1,2}(x_1, x_2) \qquad \qquad \mu_{\beta}(x_{N-1}) = \sum_{x_N} \psi_{N-1,N}(x_{N-1}, x_N)$$

• The partition function can be computed at any node:

$$Z = \sum_{x_n} \mu_{\alpha}(x_n) \mu_{\beta}(x_n)$$

• Overall, we have  $O(NK^2)$  operations to compute the marginal  $p(x_n)$ 



To compute local marginals:

- •Compute and store all forward messages,  $\mu_{\alpha}(x_n)$ .
- •Compute and store all backward messages,  $\mu_{\beta}(x_n)$
- •Compute Z once at a node  $x_m$ :

$$Z = \sum_{x_m} \mu_\alpha(x_m) \mu_\beta(x_m)$$

•Compute

$$p(x_n) = \frac{1}{Z} \mu_{\alpha}(x_n) \mu_{\beta}(x_n)$$

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for all variables required.





# Summary

- Undirected Models (also known as Markov random fields) provide a simpler method to check for conditional independence
- A MRF is defined as a factorization over clique potentials and normalized globally
- Directed models can be converted into undirected ones, but there are distributions that can be represented only in one kind of model
- For undirected Markov chains there is a very efficient inference method based on message passing

