# Constructing an UGM from a positive distribution using Local-CI

Definition: The Markov Blanket of a variable  $x_i$ ,  $MB(x_i)$  is the smallest subset of variables V that makes  $x_i$  CI of others given the Markov blanket.

$$x_i \perp V - MB(x_i) - \{x_i\} | MB(x_i)$$

#### **Theorem**

The MB of a variable is always unique for a positive distribution.

# Proof of uniqueness of MB

#### Proof.

(Not in Syllabus) We will prove by contradiction. Let  $x_i$  be a variable and  $M_1$ ,  $M_2$  be two MBs. Let  $\alpha = M_1 - M_2$  and  $\beta = M_2 - M_1$ ,

$$M = M_1 \cap M_2, W = V - (M_1 \cup M_2)$$

$$x_i \perp \!\!\!\perp V - M_2 | M_2, x_i \perp \!\!\!\perp V - M_1 | M_1,$$

This implies,  $x_i \perp \!\!\! \perp W, \alpha | M, \beta, x_i \perp \!\!\! \perp W, \beta | M, \alpha$ .

For positive distributions, this implies  $x_i \perp \!\!\! \perp W, \alpha, \beta | M$  (Intersection property of distributions Sec 2.1)

This implies that M is also a MB. But then  $M_1$ ,  $M_2$  were supposed to be minimal — a contradiction.



#### Popular undirected graphical models

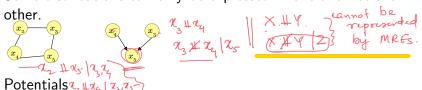
- Interacting atoms in gas and solids [ 1900]
- Markov Random Fields in vision for image segmentation
- Conditional Random Fields for information extraction
- Social networks
- Bio-informatics: annotating active sites in a protein molecules.

#### Lessons Learned

- BNs not great for representing symmetric interactions among variables. MRFs are better suited.
- Potentials are arbitrary scores not conditional probabilities
- We can draw MRFs for positive distributions by finding the Markov Blanket for each variable.
- In practice, MRFs are often constructed starting from potentials where we just connect together all variables that appear together in a potential.
- Easy to read-off all Cls using graph separability

# Comparing directed and undirected graphs

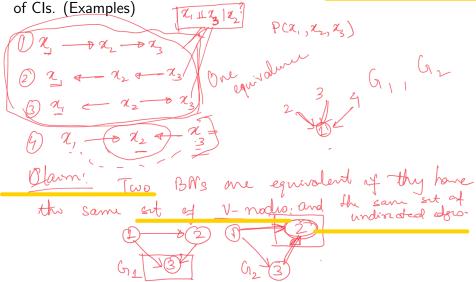
• Some distributions can only be expressed in one and not the



- Potentials 1, 11 x (7, 75)
  - Directed: conditional probabilities, more intuitive
  - Undirected: arbitrary scores, easy to set.
- Dependence structure
  - Directed: Complicated d-separation test
  - ▶ Undirected: Graph separation:  $A \perp \!\!\!\perp B \mid C$  iff C separates A and B in G.
- Often application makes the choice clear.
  - Directed: Causality -
  - Undirected: Symmetric interactions.

### Equivalent BNs

Two BN DAGs are said to be equivalent if they express the same set



#### Equivalent BNs

#### **Theorem**

Two BNs  $G_1$ ,  $G_2$  are equivalent iff they have the same skeleton and the same set of immoralities. (An immorality is a structure of the form  $x \to y \leftarrow z$  with no edge between x and z)

DOUBT - immoraity also when Y has children?



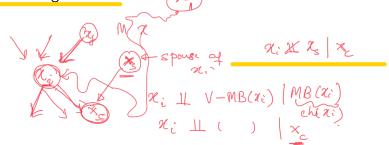
#### Converting BN to MRFs

Efficient: Using the Markov Blanket (MB) (also called the Local-CI) algorithm

The MB of a  $x_i$  in a BN can be shown to be:

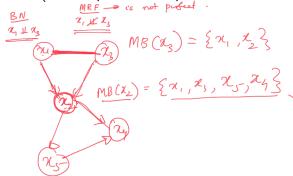
$$MB(x_i) = Pa(x_i) \cup Ch(x_i) \cup Spouse(x_i)$$

This is essentially obtained by moralizing a BN and removing all directed edges.



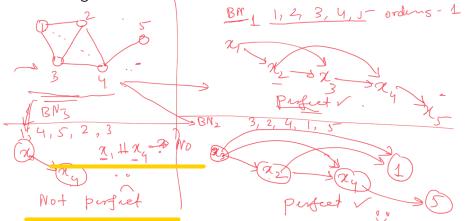
# For which BN can we create perfect MRFs?

A BN which has no immorality will not require any new edges to be added when converting to MRF. Such networks will have a perfect MRF. (Formal proof in Proposition 4.9 of KF book)



# Converting MRFs to BNs

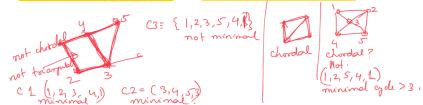
Use the BN construction algorithm (Discussed in the BN portion) starting with any random order of variables. Use the Global-Cl on the MRF to answer the conditional independence queries. Different orders will give rise to different BN.



# Which MRFs have perfect BNs

Chordal or triangulated graphs

A graph is chordal if it has no minimal cycle of length  $\geq 4$ .



#### **Theorem**

A MRF can be converted perfectly into a BN iff it is chordal.

#### Proof.

Theorems 4.11 and 4.13 of KF book

Algorithm for constructing perfect BNs from chordal MRFs to be discussed later.