

# Conditional independence and Causality

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INTELLIGENT SYSTEMS FOR PATTERN RECOGNITION (ISPR)

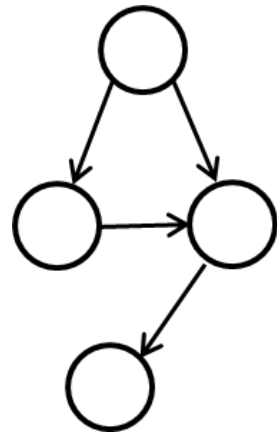
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# On the Nature of Relationships in Bayesian and Markov Networks

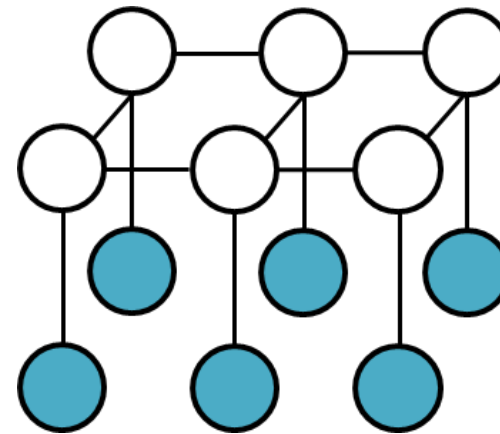
Bayesian Networks

Directed edges representing asymmetric cause-effect relationships



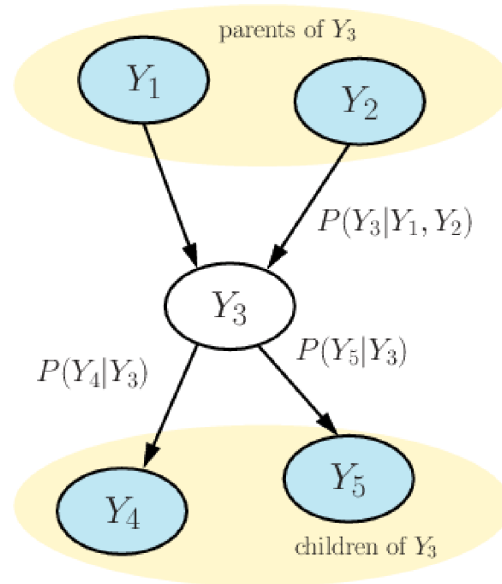
Markov Networks

Undirected edges representing symmetric relationships



*Can we reason on the structure of the graph to infer direct/indirect relationships between RVs?*

# Bayesian Network



- Directed Acyclic Graph (DAG)  $\mathcal{G} = (\mathcal{V}, \mathcal{E})$
- Nodes  $v \in \mathcal{V}$  represent random variables
  - Shaded  $\Rightarrow$  observed
  - Empty  $\Rightarrow$  un-observed
- Edges  $e \in \mathcal{E}$  describe the **conditional independence relationships**


**Conditional Probability Tables (CPT)** local to each node describe the probability distribution **given its parents**

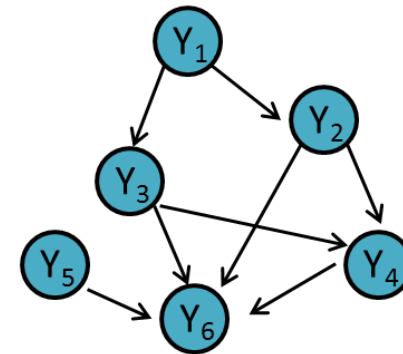
$$P(Y_1, \dots, Y_N) = \prod_{i=1}^N P(Y_i | pa(Y_i))$$

# A Simple Example

- Assume  $N$  discrete RV  $Y_i$  who can take  $k$  distinct values
- How many parameters in the **joint probability distribution**?  
 $k^N - 1$  independent parameters

How many independent parameters if **all**  $N$  variables are **independent**?  $N * (k - 1)$       What if only part of the variables are (conditionally) independent?


$$P(Y_1, \dots, Y_N) = \prod_{i=1}^N P(Y_i)$$

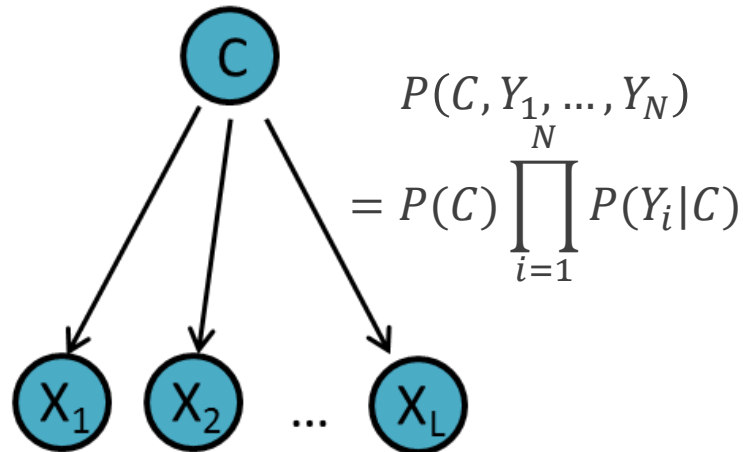


If the  $N$  nodes have a maximum of  $L$  children  $\Rightarrow (k - 1)^L \times N$  independent parameters

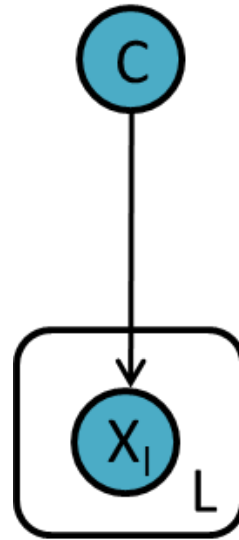


# A Compact Representation of Replication

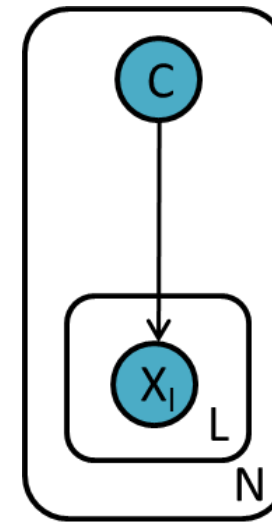
If the same **causal relationship is replicated** for a number of variables, we can compactly represent it by **plate notation**



The **Naive Bayes Classifier**

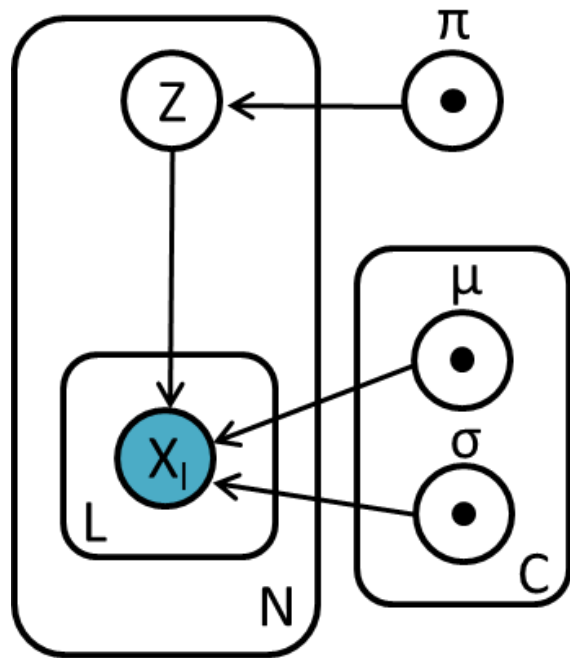


Replication for  $L$  attributes



Replication for  $N$  data samples

# Full Plate Notation



Gaussian Mixture Model

- Boxes denote **replication** for a number of times denoted by the **letter in the corner**
- Shaded nodes are **observed** variables
- Empty nodes denote un-observed **latent** variables
- Black seeds (optional) identify **model parameters**
  - $\pi \rightarrow$  multinomial prior distribution
  - $\mu \rightarrow$  means of the  $C$  Gaussians
  - $\sigma \rightarrow$  std of the  $C$  Gaussians

# Local Markov Property

## Definition (Local Markov property)

Each node / random variable is conditionally independent of **all its non-descendants** given a **joint state of its parents**

$$Y_v \perp Y_{V \setminus \text{ch}(v)} \mid Y_{\text{pa}(v)} \text{ for all } v \in V$$

*Party* and *Study* are **marginally** independent

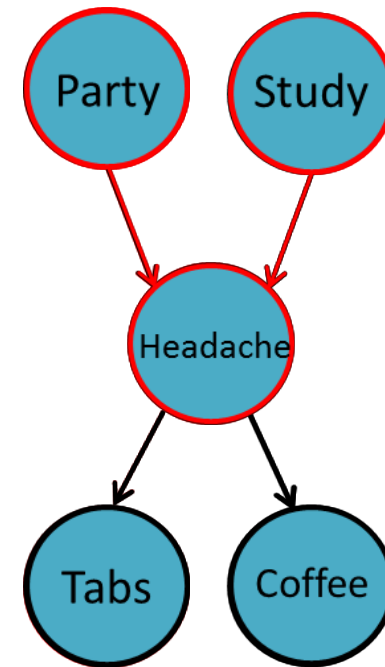
- *Party*  $\perp$  *Study*

However, local Markov property **does not support**

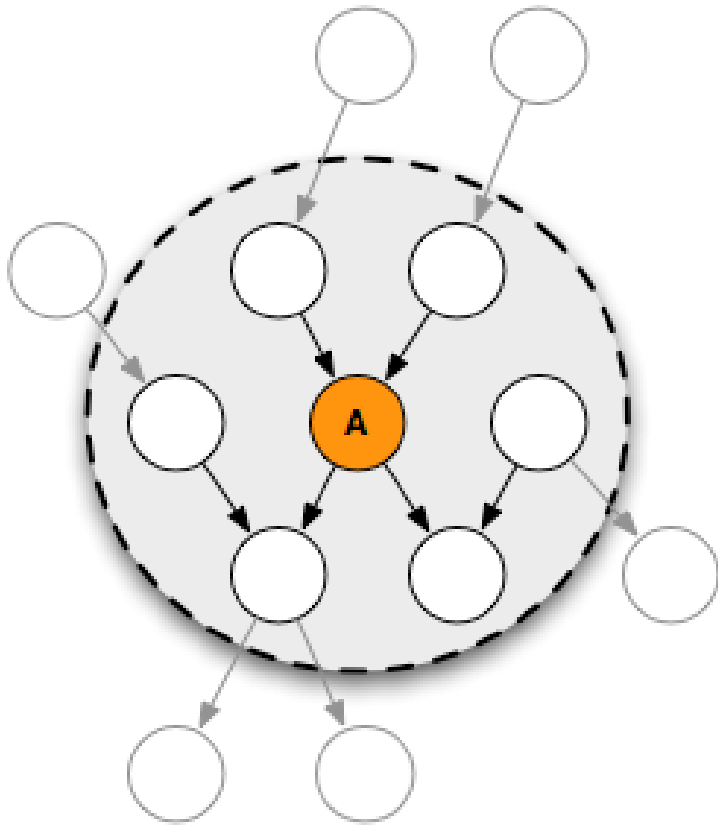
- *Party*  $\perp$  *Study* | *Headache*

- *Tabs*  $\perp$  *Party*

But *Party* and *Tabs* are **independent given Headache**



# Markov Blanket



- The **Markov Blanket**  $Mb(A)$  of a node  $A$  is the minimal set of vertices that **shield the node** from the rest of Bayesian Network
- The behavior of a node can be **completely determined and predicted** from the knowledge of its Markov blanket

$$P(A|Mb(A), Z) = P(A|Mb(A)) \quad \forall Z \notin Mb(A)$$

- The Markov blanket of  $A$  contains
  - Its parents  $pa(A)$
  - Its children  $ch(A)$
  - Its children's parents  $pa(ch(A))$

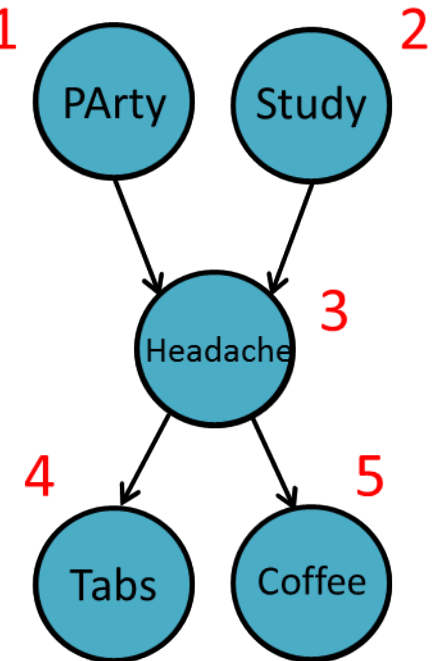




# Joint Probability Factorization

An application of **Chain rule** and **Local Markov Property** <sup>1</sup>

1. Pick a **topological ordering** of nodes
2. Apply **chain rule** following the order
3. Use the **conditional independence assumptions**



$$\begin{aligned} P(PA, S, H, T, C) &= \\ &P(PA) \cdot P(S|PA) \cdot P(H|S, PA) \cdot P(T|H, S, PA) \cdot P(C|T, H, S, PA) \\ &= P(PA) \cdot P(S) \cdot P(H|S, PA) \cdot P(T|H) \cdot P(C|H) \end{aligned}$$



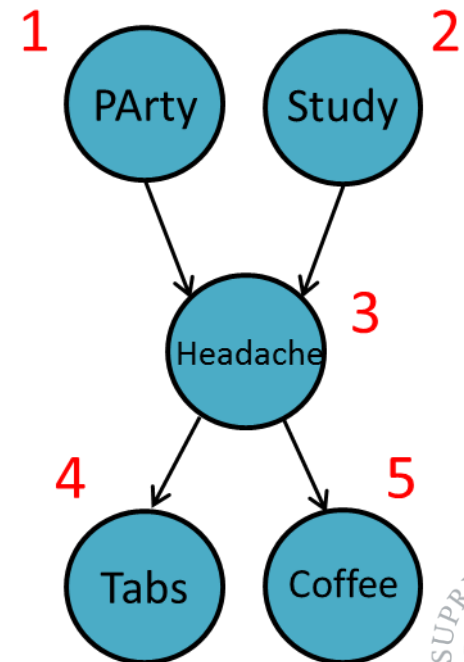
# (Ancestral) Sampling of a BN

A BN describes a generative process for observations

1. Pick a **topological ordering** of nodes
2. Generate data by **sampling from the local conditional probabilities** following this order

Generate  $i$ -th sample for each variable  $PA, S, H, T, C$

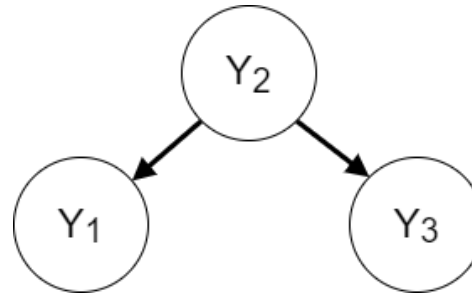
1.  $pa_i \sim P(PA)$
2.  $s_i \sim P(S)$
3.  $h_i \sim P(H|S = s_i, PA = pa_i)$
4.  $t_i \sim P(T|H = h_i)$
5.  $c_i \sim P(C|H = h_i)$



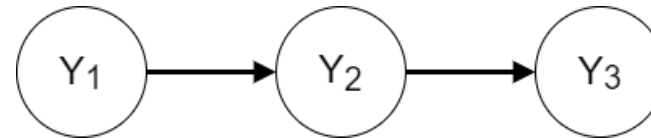
# Fundamental BN structures

There exist 3 fundamental substructures that determine the conditional independence relationships in a Bayesian network

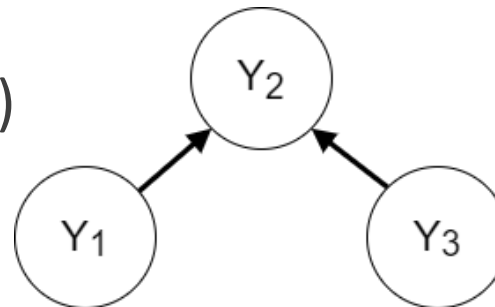
- Tail to tail (Common Cause)



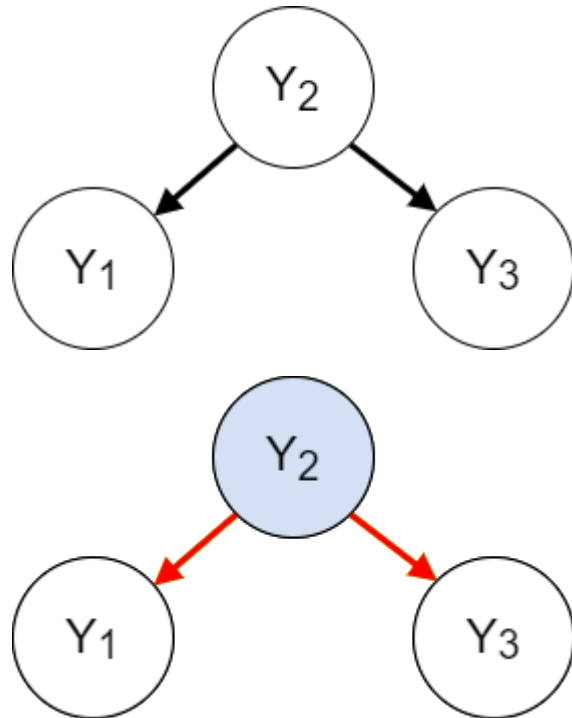
- Head to tail (Causal Effect)



- Head to head (Common Effect)



# Tail to Tail Connections



- Corresponds to
$$P(Y_1, Y_3|Y_2)P(Y_2) = P(Y_1|Y_2)P(Y_3|Y_2)P(Y_2)$$
- If  $Y_2$  is unobserved then  $Y_1$  and  $Y_3$  are marginally dependent

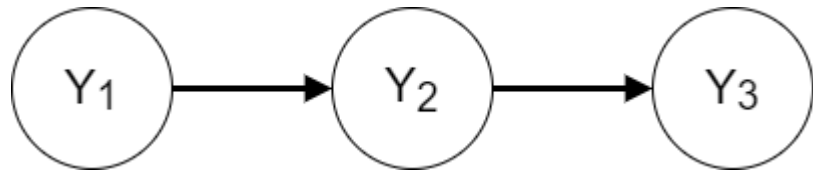
$$Y_1 \not\perp Y_3$$

- If  $Y_2$  is observed then  $Y_1$  and  $Y_3$  are conditionally independent

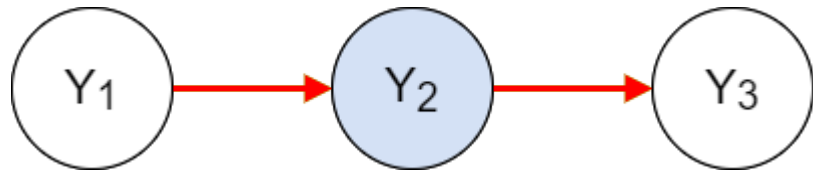
$$Y_1 \perp Y_3|Y_2$$

When  $Y_2$  is observed is said to **block the path** from  $Y_1$  to  $Y_3$

# Head to Tail Connections



- Corresponds to
$$P(Y_1, Y_2, Y_3) = P(Y_1)P(Y_2|Y_1)P(Y_3|Y_2)$$
$$= P(Y_1|Y_2)P(Y_3|Y_2)P(Y_2)$$



Observed  $Y_2$  blocks the path from  $Y_1$  to  $Y_3$

- If  $Y_2$  is unobserved then  $Y_1$  and  $Y_3$  are marginally dependent

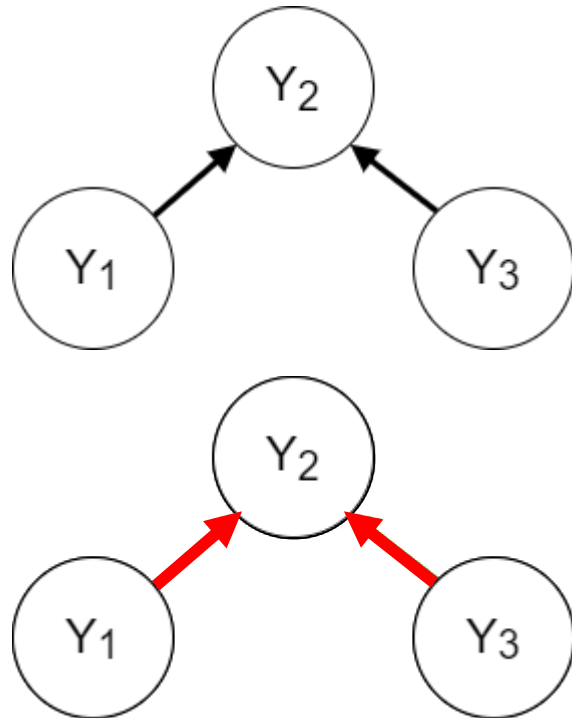
$$Y_1 \not\perp Y_3$$

- If  $Y_2$  is observed then  $Y_1$  and  $Y_3$  are conditionally independent

$$Y_1 \perp Y_3 | Y_2$$



# Head to Head Connections



- Corresponds to

$$P(Y_1, Y_2, Y_3) = P(Y_1)P(Y_3)P(Y_2|Y_1, Y_3)$$

- If  $Y_2$  is observed then  $Y_1$  and  $Y_3$  are conditionally dependent

$$Y_1 \not\perp Y_3 | Y_2$$

- If  $Y_2$  is unobserved then  $Y_1$  and  $Y_3$  are marginally independent

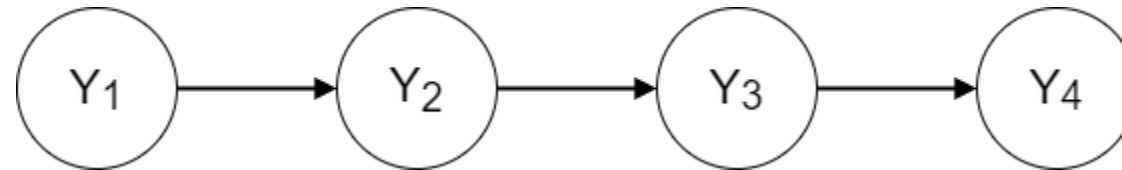
$$Y_1 \perp Y_3$$

If any  $Y_2$  descendant is observed it unlocks the path

# Derived Conditional Independence Relationships

A Bayesian Network represents the local relationships encoded by the 3 basic structures plus the **derived relationships**

Consider



Local Markov Relationships

$$Y_1 \perp Y_3 | Y_2$$

$$Y_4 \perp Y_1, Y_2 | Y_3$$

Derived Relationship

$$Y_1 \perp Y_4 | Y_2$$



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# d-Separation

## Definition (d-separation)

Let  $r = Y_1 \leftrightarrow \dots \leftrightarrow Y_2$  be an **undirected path** between  $Y_1$  and  $Y_2$ , then  $r$  is **d-separated by  $Z$**  if there exist at least one node  $Y_c \in Z$  for which path  $r$  is blocked.

In other words, **d-separation** holds if at least one of the following holds

- $r$  contains an **head-to-tail** structure  $Y_i \rightarrow Y_c \rightarrow Y_j$  (or  $Y_i \leftarrow Y_c \leftarrow Y_j$ ) and  $Y_c \in Z$
- $r$  contains a **tail-to-tail** structure  $Y_i \leftarrow Y_c \rightarrow Y_j$  and  $Y_c \in Z$
- $r$  contains an **head-to-head** structure  $Y_i \rightarrow Y_c \leftarrow Y_j$  and **neither  $Y_c$  nor its descendants are in  $Z$**





# Markov Blanket and d-Separation

## Definition (Nodes d-separation)

Two nodes  $Y_i$  and  $Y_j$  in a BN  $\mathcal{G}$  are said to be **d-separated by**  $Z \subset \mathcal{V}$  (denoted by  $Dsep_{\mathcal{G}}(Y_i, Y_j | Z)$ ) if and only if all undirected paths between  $Y_i$  and  $Y_j$  are d-separated by  $Z$

## Definition (Markov Blanket)

The Markov blanket  $Mb(Y)$  is the minimal set of nodes which d-separates a node  $Y$  from all other nodes (i.e. it makes  $Y$  conditionally independent of all other nodes in the BN)

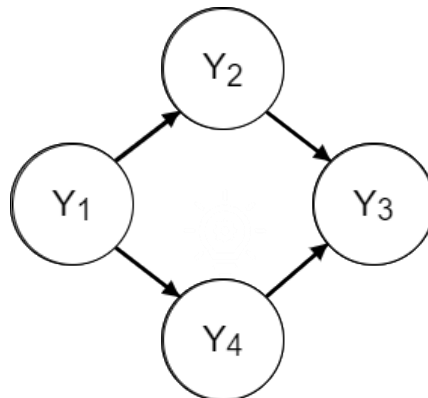
$$Mb(Y) = \{pa(Y), ch(Y), pa(ch(Y))\}$$



# Are Directed Models Enough?

- Bayesian Networks are used to model **asymmetric dependencies** (e.g. causal)
- What if we want to model **symmetric dependencies**
  - Bidirectional effects, e.g. spatial dependencies
  - Need **undirected** approaches

Directed models cannot represent some (bidirectional) dependencies in the distributions

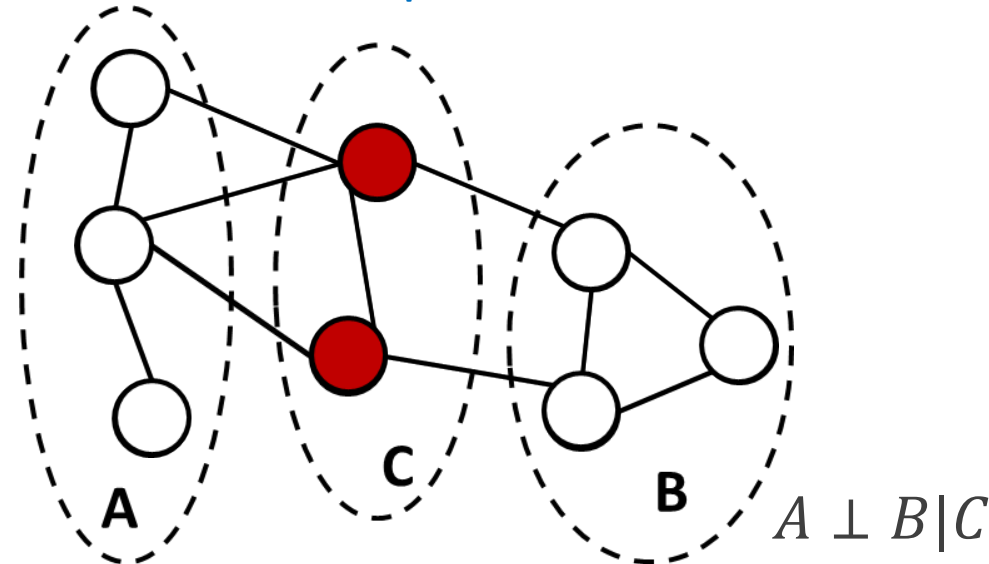


What if we want to represent  $Y_1 \perp Y_3 | Y_2, Y_4$ ?  
What if we also want  $Y_2 \perp Y_4 | Y_1, Y_3$ ?

Cannot be done in BN! Need undirected model

# Markov Random Fields

What is the **undirected equivalent** of **d-separation** in directed models?



Again it is based on node separation, although it is way simpler!

- Node subsets  $A, B \subset \mathcal{V}$  are **conditionally independent** given  $C \subset \mathcal{V} \setminus \{A, B\}$  if all paths between nodes in  $A$  and  $B$  pass through at least one of the nodes in  $C$
- The **Markov Blanket** of a node includes all and only its **neighbors**

# Joint Probability Factorization

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What is the **undirected equivalent** of **conditional probability factorization** in directed models?

- We seek a **product of functions** defined over a set of nodes associated with some **local property** of the graph
- Markov blanket tells that **nodes that are not neighbors are conditionally independent** given the remainder of the nodes

$$P(X_v, X_i | X_{\mathcal{V} \setminus \{v, i\}}) = P(X_v | X_{\mathcal{V} \setminus \{v, i\}})P(X_i | X_{\mathcal{V} \setminus \{v, i\}})$$

- Factorization should be chosen in such a way that nodes  $X_v$  and  $X_i$  are not in the same factor

**What is a well-known graph structure that includes only nodes that are pairwise connected?**



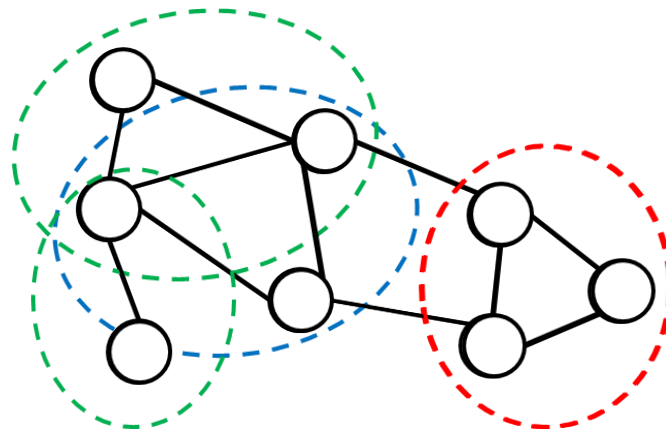
# Cliques

## Definition (Clique)

A subset of nodes  $C$  in graph  $G$  such that  $G$  contains an edge between all pair of nodes in  $C$

## Definition (Maximal Clique)

A clique  $C$  that cannot include any further node from the graph without ceasing to be a clique



# Maximal Clique Factorization

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Define  $\mathbf{X} = X_1, \dots, X_N$  as the RVs associated to the  $N$  nodes in the undirected graph  $\mathcal{G}$

$$P(\mathbf{X}) = \frac{1}{Z} \prod_C \psi(\mathbf{X}_C)$$

- $\mathbf{X}_C \rightarrow$  RV associated with nodes in the maximal clique  $C$
- $\psi(\mathbf{X}_C) \rightarrow$  potential function over the maximal cliques  $C$
- $Z \rightarrow$  partition function ensuring normalization

$$Z = \sum_{\mathbf{X}} \prod_C \psi(\mathbf{X}_C)$$

Partition function is the **computational bottleneck** of undirected modes:  
e.g.  $O(K^N)$  for  $N$  discrete RV with  $K$  distinct values

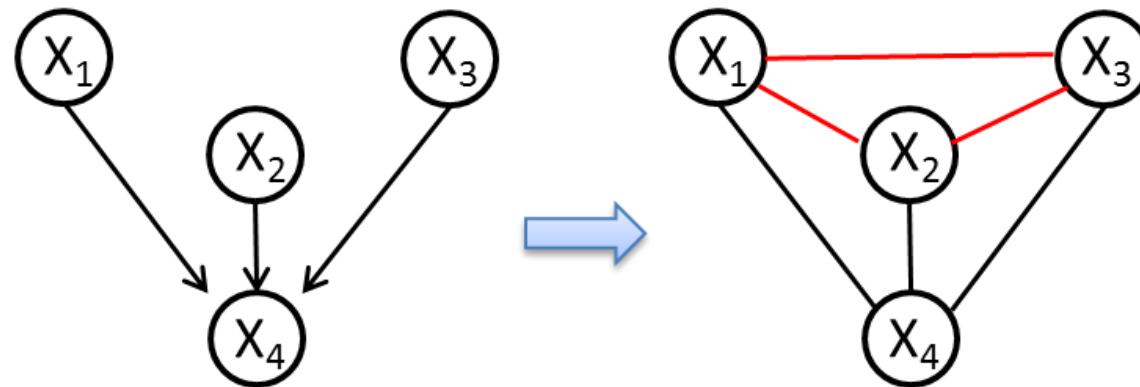


# From Directed To Undirected

Straightforward in some cases



Requires a little bit of thinking for **v-structures**

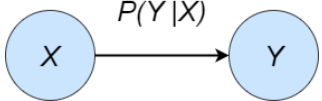
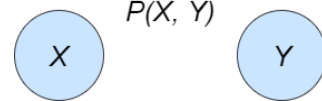


**Moralization** a.k.a. marrying of the parents

# Learning Causation (from data)

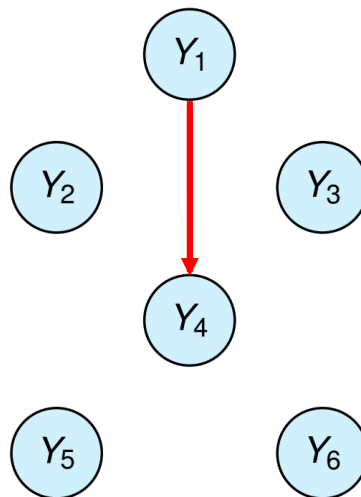


# Learning with Bayesian Networks

		Structure	
		Fixed Structure	Fixed Variables
Data	Complete	 <p>Naive Bayes Calculate Frequencies (ML)</p>	 <p>Discover dependencies from the data Structure Search Independence tests</p>
	Incomplete	<p>Latent variables EM Algorithm (ML) MCMC, VBEM (Bayesian)</p> <p><b>Parameter Learning</b></p>	<p>Difficult Problem Structural EM</p> <p><b>Structure Learning</b></p>

# The Structure Learning Problem

$Y_1$	$Y_2$	$Y_3$	$Y_4$	$Y_5$	$Y_6$
1	2	1	0	3	4
4	0	0	0	1	2
...	...	...	...	...	...
...	...	...	...	...	...
0	0	1	3	2	1



- Observations are given for a set of **fixed random variables**
- Network structure is not specified
  - Determine which arcs exist in the network (**causal relationships**)
  - Compute Bayesian network parameters (**conditional probability tables**)
- Determining causal relationships between variables entails
  - Deciding on **arc presence**
  - **Directing edges**



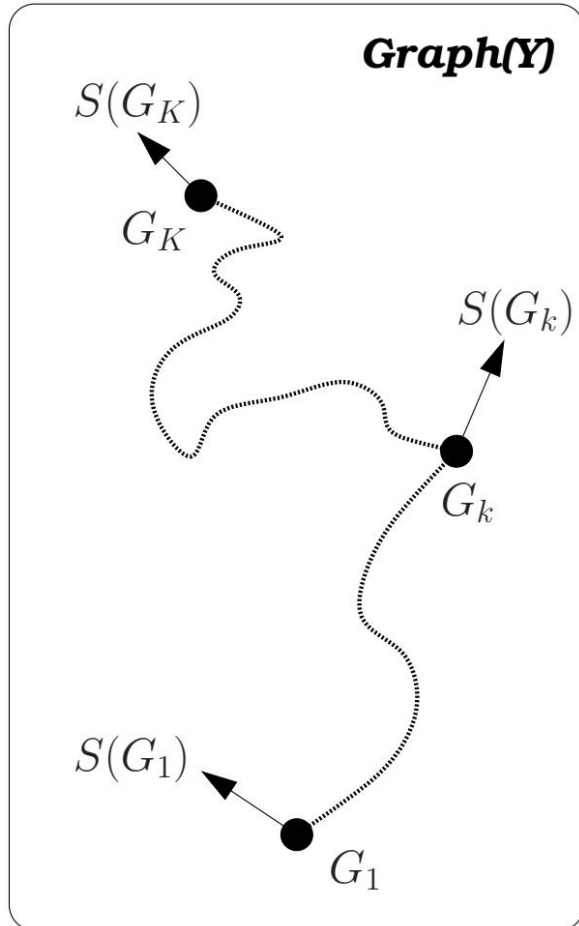
# Structure Finding Approaches

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- Search and Score
  - Model selection approach
  - Search in the space of the graphs
- Constraint Based
  - Use tests of conditional independence
  - Constrain the network
- Hybrid
  - Model selection of constrained structures



# Search & Score



- Search the space  $Graph(\mathbf{Y})$  of graphs  $G_k$  that can be built on the random variables  $\mathbf{Y} = Y_1, \dots, Y_N$
- Score each structure by  $S(G_k)$
- Return the highest scoring graph  $G^*$
- Two fundamental aspects
  - Scoring function
  - Search strategy

# Scoring Function

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- Fundamental properties
  - **Consistency** - Same score for graphs in the same equivalence class
  - **Decomposability** - Can be locally computed
- Approaches
  - **Information theoretic** - Based on data likelihood plus some model-complexity penalization terms (AIC, BIC, MDL, ...)
  - **Bayesian** – Score the structures using a graph posterior (likelihood + proper prior choice)

$$\log P(D|G) \approx \sum_D \sum_X \log \tilde{P}(x|\mathbf{pa}(x)) + \log P(G)$$



# Search Strategy

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- Finding maximal scoring structures is NP complete (Chickering, 2002)
- **Constrain search strategy**
  - Starting from a candidate structure **modify iteratively by local operations** (edge/node addition or deletion)
  - Each operation has a cost
  - **Cost optimization** problem: greedy hill-climbing, simulated annealing, ...
- **Constrain search space**
  - **Known node order** – Can reduce the search space to the parents of each node (Markov Blanket)
  - Search in the space of **structure equivalence classes** (GES algorithm)
  - Search in the space of **node orderings** (Friedman and Koller, 2003)



# Constraint-based Models

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- Tests of **conditional independence**  $I(X_i, X_j|Z)$  determine edge presence (**network skeleton**)
- Based on measures of association between two variables/nodes  $X_i$  and  $X_j$ , given their neighbor nodes  $Z$ 
  - Conditional **mutual information**
  - Statistical **hypothesis testing** on association measures with a **known distribution**, e.g.  $\chi^2$

$$G^2(X_i, X_j|Z) = 2 \sum_{x_i, x_j, z} n_D(x_i, x_j, z) \frac{n_D(x_i, x_j, z) n_D(z)}{n_D(x_i, z) n_D(x_j, z)}$$

- Use deterministic rules based on local Markovian dependencies to determine edge orientation (**DAG**)



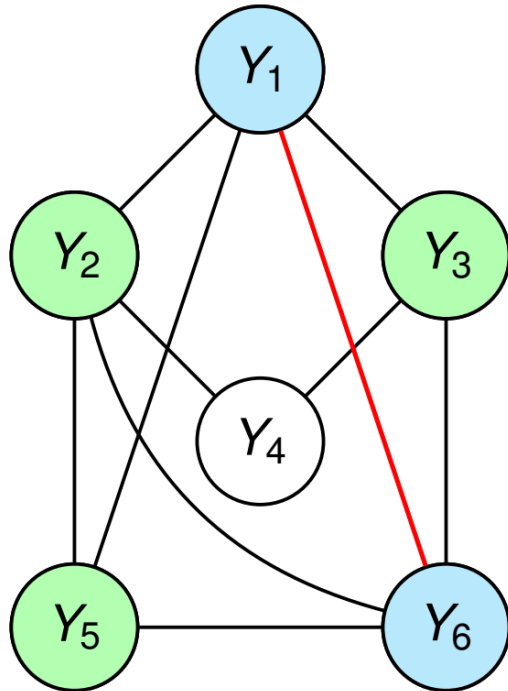
# Testing Strategy

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- Choice of the **testing order** is fundamental for avoiding a **super-exponential** complexity
- Level-wise testing
  - Tests  $I(X_i, X_j|Z)$  are performed in order of **increasing size** of the conditioning set  $Z$  (starting from empty  $Z$ )
  - PC algorithm (Spirtes, 1995)
- Node-wise testing
  - Tests are performed on a **single edge** at the time, exhausting independence checks on **all conditioning variables**
  - TPDA Algorithm
- Nodes that enter  $Z$  are chosen in the **neighborhood** of  $X_i$  and  $X_j$



# PC Algorithm



Initialize a fully connected graph  $\mathcal{G} = (\mathcal{V}, \mathcal{E})$

**for each** edge  $(Y_i, Y_j) \in \mathcal{V}$

- **if**  $I(Y_i, Y_j)$  **then** prune  $(Y_i, Y_j)$

$K \leftarrow 1$

**for each** test of order  $K = |\mathcal{Z}|$

- **for each** edge  $(Y_i, Y_j) \in \mathcal{V}$

- $\mathcal{Z} \leftarrow$  set of conditioning sets of  $K$ -th order for  $Y_i, Y_j$

- **if**  $I(Y_i, Y_j|z)$  **for any**  $z \in \mathcal{Z}$  **then** prune  $(Y_i, Y_j)$

- $K \leftarrow K + 1$

**return**  $\mathcal{G}$



# Hybrid Models

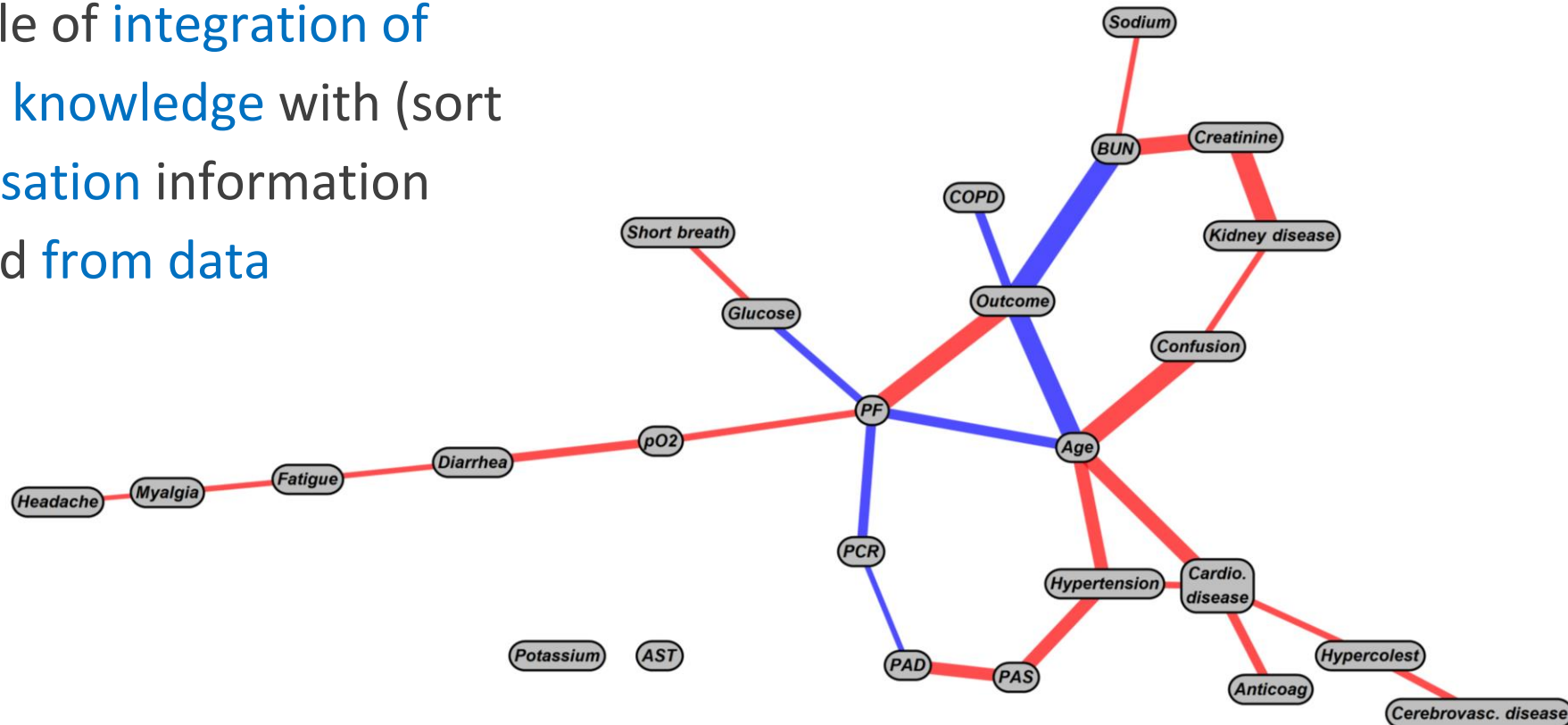
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- Multi-stage algorithms combining previous approaches
- Independence tests to find a sub-optimal skeleton (**good starting point**)
- Search and score **starting from the skeleton**
  - Skeleton refinement
  - Edge orientation
- **Max-Min Hill Climbing** (MMHC) model
  - Optimized constraint-based approach to reconstruct the skeleton (**Max-Min Parents and Children**)
  - Use the **candidate parents** in the skeleton to run a search and score approach



# Learning a COVID-19 causal model

Example of integration of clinical knowledge with (sort of) causation information inferred from data



# Take Home Messages

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- Directed graphical models
  - Represent **asymmetric (causal) relationships** between RV and conditional probabilities in compact way
  - Difficult to assess conditional independence (v-structures)
  - Ok for **prior knowledge** and **interpretation**
- Undirected graphical models
  - Represent **bi-directional relationships** (e.g. constraints)
  - Factorization in terms of generic **potential functions** (not probabilities)
  - Easy to assess conditional independence, but **difficult to interpret**
  - Serious **computational issues** due to normalization factor
- Structure learning to **infer multivariate causation relationships** from data



# Next Two Lectures

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## Hidden Markov Model (HMM)

- A dynamic graphical model for sequences
- Unfolding learning models on structures
- Exact inference on a chain with observed and unobserved variables
- The Expectation-Maximization algorithm for HMMs

