

Learning in Bayesian Networks



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Berlin: 20.06.2002

Overview

1. Bayesian Networks

- Stochastic Networks
- Bayesian statistics
- Use of Bayesian Networks
 - Inference given a network,
 - Learn the network from data,
 - Causal Inference

2. Analysis of Gene Expression Data

- Application: Cell Cycle Expression Patterns



References

1. Bayesian Networks

HECKERMAN, David: *A Tutorial on Learning with Bayesian Networks*, MSR-TR-95-06

2. Analysis of Gene Expression Data

FRIEDMAN *et. al.*: *Using Bayesian Networks to Analyze Expression Data*, RECOMB 1999.

SPELLMAN *et. al.*: *Comprehensive identification of cell cycle-regulated genes of the yeast *sac. cer.* by microarray hybridization*, *Molecular Biology of the Cell*, 9:3273–3297, 1998.



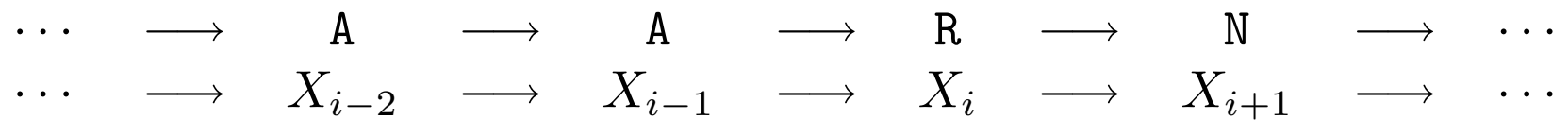
Graphical Models

1. Markov Chains
 2. Hidden Markov Models
 3. Markov Random Fields
 4. Bayesian Networks
- ...



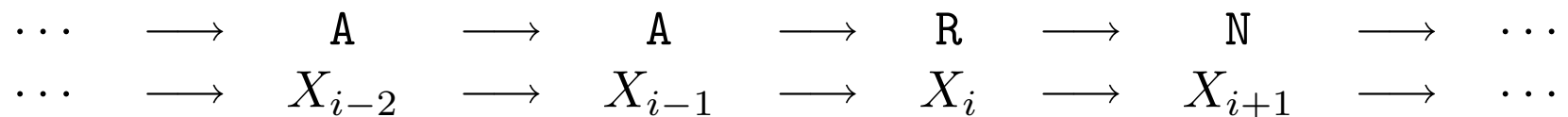
Markov Chains

Example: Evolution of a protein sequence modelled by a first-order Markov Chain.



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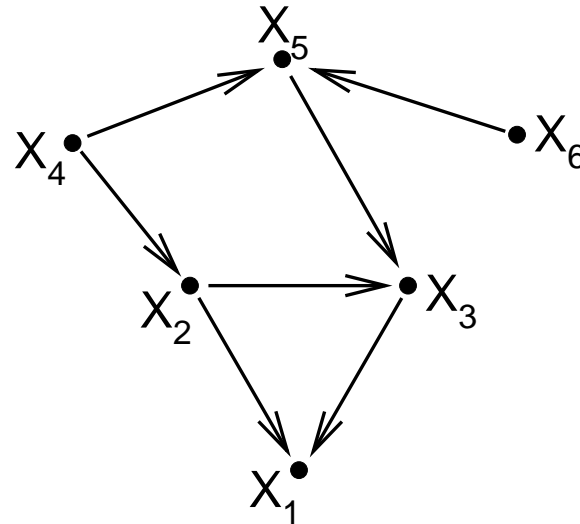
- **Markov Condition:**

$$P(X_i | X_{i-1}, X_{i-2}, \dots) = P(X_i | X_{i-1})$$

- **Local probability** in X_i depends only on direct predecessor X_{i-1} .
- **Conditional Independence:** Given its predecessor, X_i is independent from the other nodes in the graph.



A more complex example



- **Markov Condition:** Each variable X_i is independent of its non-descendants given its parents.
- **Local probability** in X_i depends only on the parents.
- **Conditional Independence:** Given its parents, X_i is independent from the other nodes in the graph.



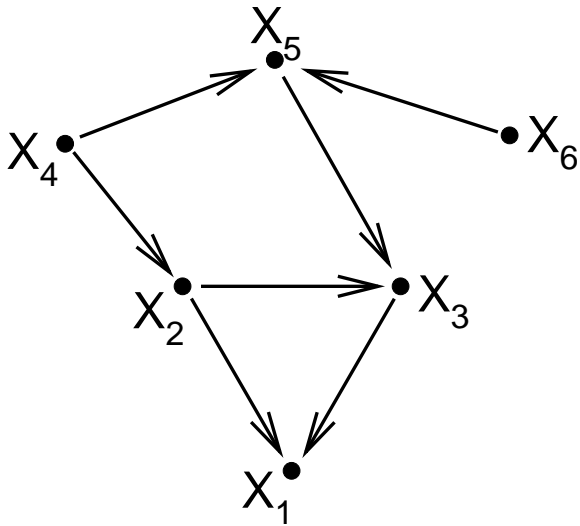
Stochastic Networks I

A **Bayesian Network** for $\mathbf{X} = \{X_1, \dots, X_n\}$ consists of

- a **network structure** \mathcal{S}
 - directed acyclic graph (DAG),
 - nodes \leftrightarrow variables,
 - lack of arc \leftrightarrow conditional independence
- a set of **probability distributions** \mathcal{P}
 - locally: conditional distribution of a variable given its parents in \mathcal{S} :
$$\mathcal{P} = \{ P(X_i \mid pa_i) \}$$



Stochastic Networks II



$(\mathcal{S}, \mathcal{P})$ encode the joint distribution

$$P(\mathbf{X}) = \prod_{i=1}^n P(X_i \mid pa_i)$$



Equivalence of Networks

Markov equivalence

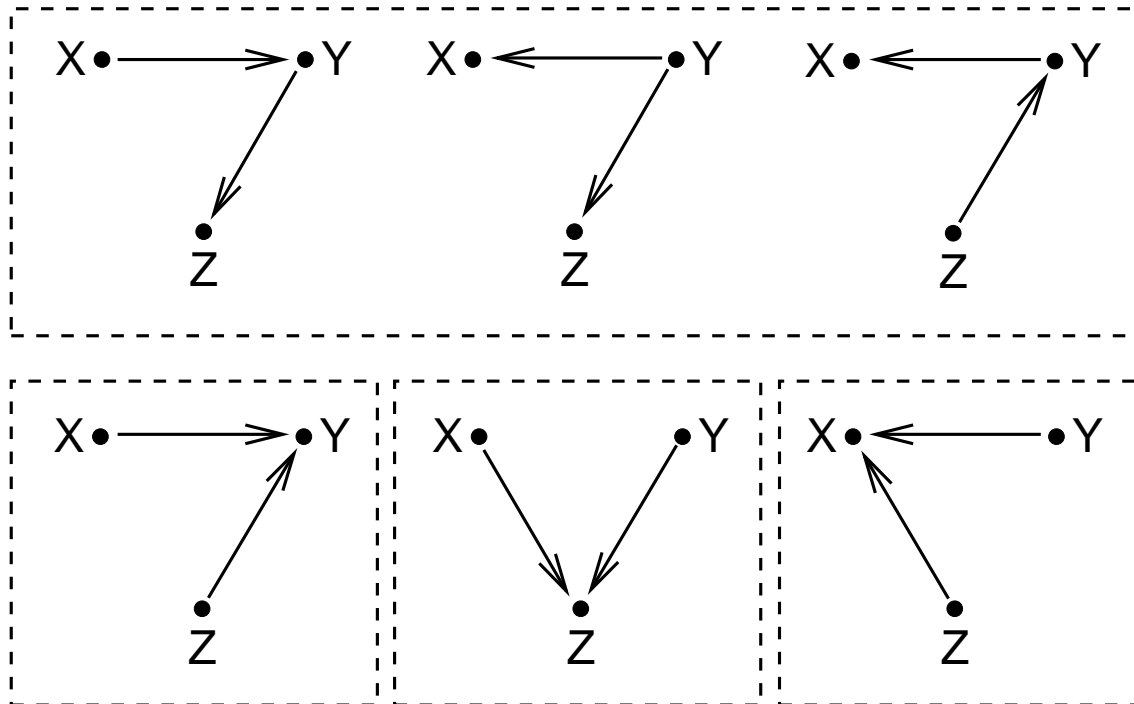
$\mathcal{S}_1 \stackrel{M}{\sim} \mathcal{S}_2$ if both structures represent the same set of independence assertions.



Equivalence of Networks

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Representation of an equivalence class

Theorem 1: [Verma and Pearl, 1990] Two DAGs are equivalent iff they have *the same skeletons and the same v-structures*.



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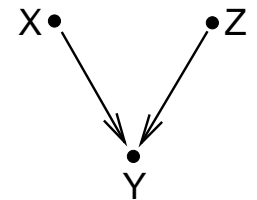
- **skeleton:**

the undirected graph resulting from ignoring the directionality of every edge.

- **v-structure:**

an ordered triple of nodes (X, Y, Z) in a graph such that

- (1) $X \longrightarrow Y$ and $Z \longrightarrow Y$
- (2) X and Z are not adjacent.



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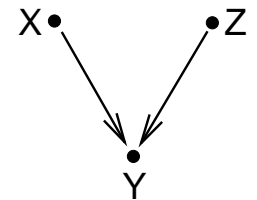
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Using Theorem 1 we can uniquely represent a equivalence class of DAGs by a **partially directed graph** (PDAG).



DAG-2-PDAG

Construction:

The PDAG identifying the equivalence class of a given DAG contains

- a **directed edge** for every edge participating in a v-structure, and
- an **undirected edge** for all other edges.

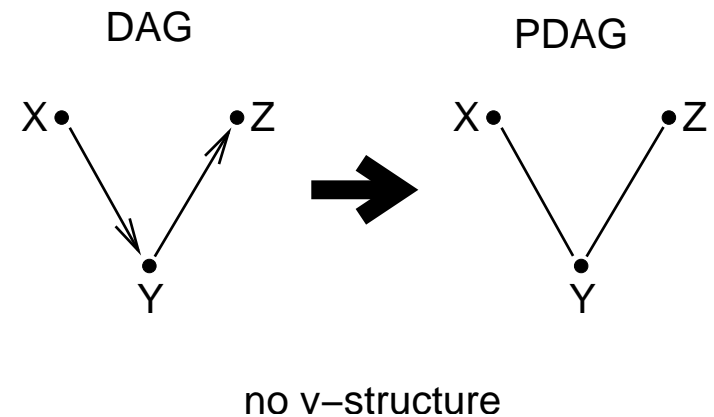
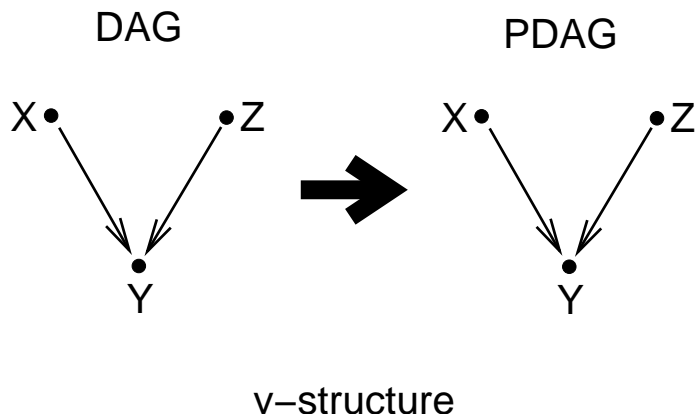


DAG-2-PDAG

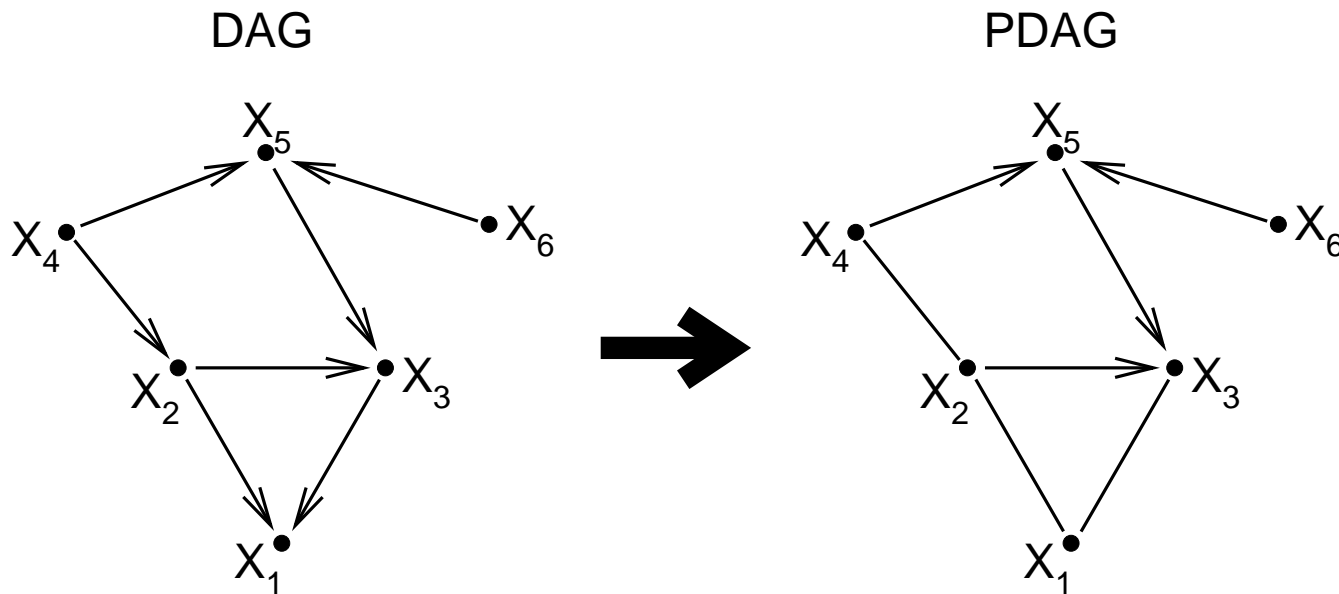
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again: a more complex example



- $X \rightarrow Y$: all members of the class agree in this arc;
- $X - Y$: some $X \rightarrow Y$ and some $Y \rightarrow X$.



What is Probability?

Frequentist Answer:

Probability is ...

- a physical property of the world
- measured by repeated trials

Bayesian Answer:

Probability is ...

- a personal degree of belief
- arbitrary (techniques for sensible choice)



Bayes Formula

Bayes Formula:

$$P(\vartheta \mid D) = \frac{P(\vartheta) P(D \mid \vartheta)}{P(D)}$$

Joint distribution:

$$P(\vartheta, D) = P(\vartheta) P(D \mid \vartheta)$$



Bayes Formula

Bayes Formula:

$$\underbrace{P(\vartheta | D)}_{\text{Posterior}} = \frac{\overbrace{P(\vartheta)}^{\text{Prior}} \overbrace{P(D | \vartheta)}^{\text{likelihood}}}{P(D)}$$

Joint distribution:

$$P(\vartheta, D) = P(\vartheta) P(D | \vartheta)$$

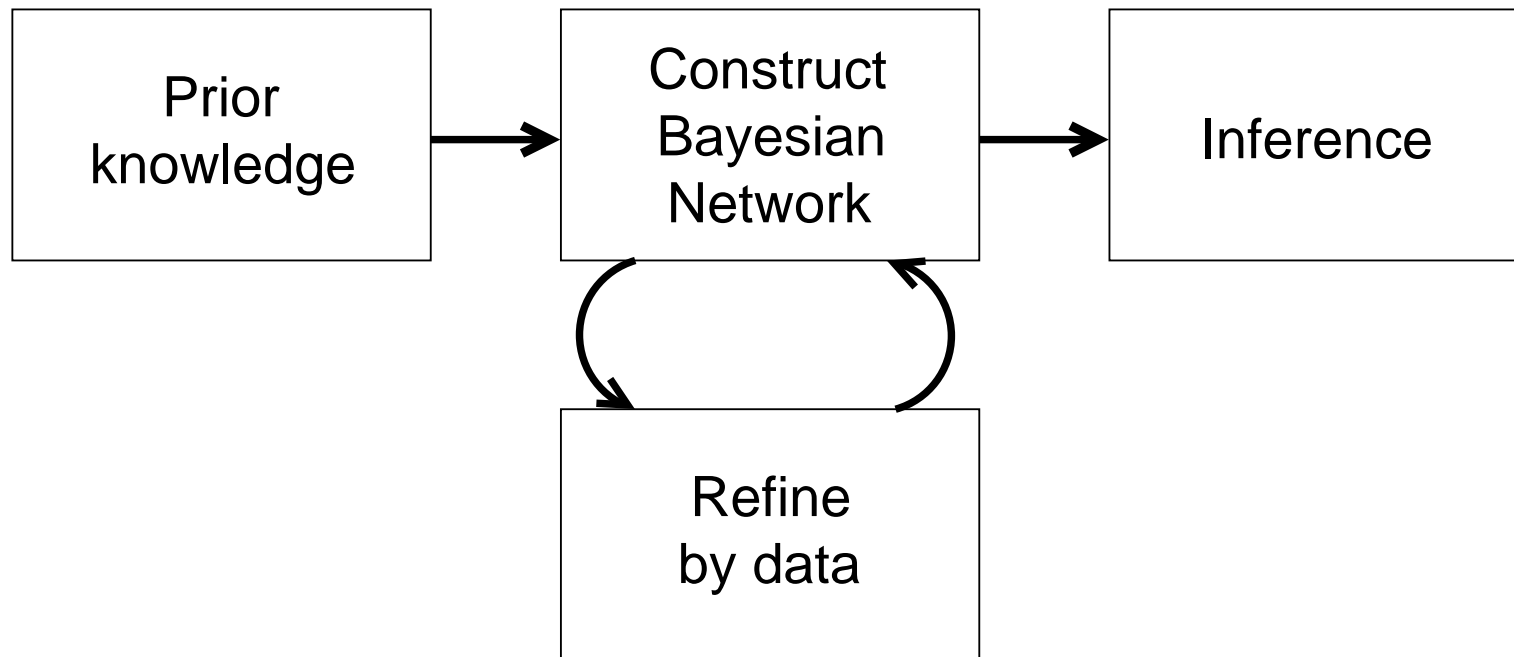


How to use a Bayesian Network?

1. **Probabilistic inference:**
Determine probability of interest from model
2. **Learn Network:**
Infer Structure and Probabilities from Data
3. **Causal Inference:**
Detect causal patterns.



1. Inference in a Bayesian Network



1.1 Construction

- Determine the **variables to model**,
- **build DAG** that encodes conditional independence
edge: cause \longrightarrow effect
- assess **local probability distributions** $P(X_i \mid pa_i)$

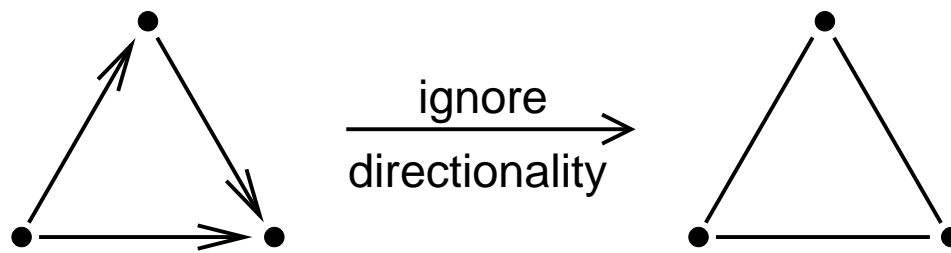


1.2 Probabilistic Inference

Probabilistic Inference: compute a probability of interest given a model.

⇒ Use Bayes theorem and simplify by conditional independence

Source of difficulty: undirected cycles



1.3 Refinement

Using data to update the probabilities of a given Bayesian Network structure:

$$P(\mathbf{X} \mid \vartheta, \mathcal{S}) = \prod_{i=1}^n P(X_i \mid pa_i, \vartheta_i, \mathcal{S})$$

Uncertainty about local distributions is encoded in Prior $P(\vartheta \mid \mathcal{S})$

Refinement: Compute Posterior from Prior.

$$\text{Prior } P(\vartheta \mid \mathcal{S}) \implies P(\vartheta \mid D, \mathcal{S}) \text{ Posterior}$$



2. Learning a network structure

Learning: find network structure which fits the prior knowledge and data.

- **Measure** this by a score function, e. g.

$$\text{Score}_D(\mathcal{S}) = \log \underbrace{p(\mathcal{S})}_{\text{Prior}} + \log \underbrace{p(D|\mathcal{S})}_{\text{likelihood}}$$

- **Search** for highscoring structure by *greedy search*, *simulated annealing*, ...



3. Learning Causal Relationships

Objective: Determine the “flow of causality” in a system.

Causal Graph: DAG \mathcal{C} is a *causal graph* for $\mathbf{X} = \{X_1, \dots, X_n\}$ iff

nodes n_i \Leftrightarrow variables X_i ,

$n_i \rightarrow n_j$ \Leftrightarrow X_i is **direct cause** of X_j .



Causal Networks vs. Bayesian Networks

1. Connection

Causal Markov Condition:

Given the values of a variables immediate causes,
it is independent of its earlier causes.

Thus:

\mathcal{C} is a causal graph for $\mathbf{X} \implies$

\mathcal{C} is a Bayesian Network structure for the joint probability distribution of \mathbf{X} .

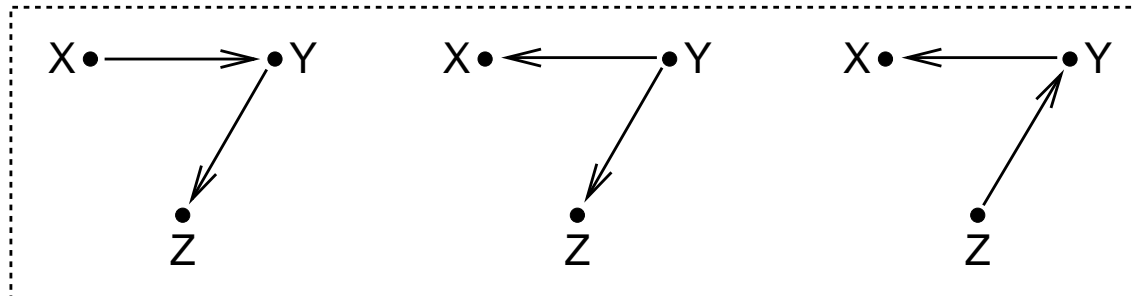


Causal Networks vs. Bayesian Networks

2. Differences

A **Bayesian Network** models the distribution of observations.

A **Causal Network** models the distribution of observations *and* effects of interventions.

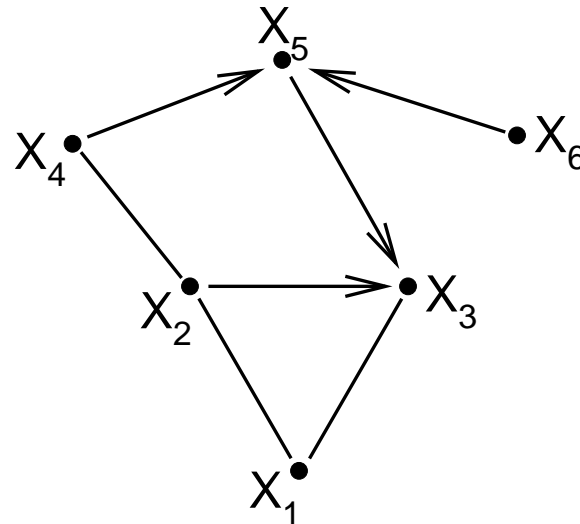


Equivalent as Bayesian Networks, **but not** equivalent as Causal Networks.



Causal Inference from Data

From observations alone, we can only learn a PDAG: a whole equivalence class of structures.



$X \longrightarrow Y$ in the PDAG: all networks agree on this directed arc.
 \implies Infer causal direction **X causes Y**.



Application: Cell Cycle in yeast

- Data from SPELLMAN *et. al.*: *Comprehensive identification of cell cycle-regulated genes of the yeast sac. cer. by microarray hybridization*, *Molecular Biology of the Cell*, 9:3273–3297, 1998.
- contains 76 samples of all the yeast genome (time series in few minutes intervals)
- Spellman *et. al.* identified 800 cell cycle regulated genes, and clustered them: 250 genes in 8 clusters
- Friedman *et. al.* analysed these 250 genes by a bayesian network.



Data Representation

Random Variables:

- Expression levels of each gene
- In addition:
 - experimental conditions
 - temporal indication (cell cycle phase)
 - background variables
 - exogenous cellular conditions



Discretization

Focus on **qualitative aspects** of the data.

Discretize gene expression data in three categories:

- 1 lower than control
- 0 equal
- 1 greater



Pairwise Features

Small datasets with many variables: many different networks are reasonable explanations of the data

⇒ focus on features that are common to most of these networks

1. Markov relations:

Is X an direct relative of Y ?
(local property)

2. Order relations:

Is X an ancestor of Y in all networks of a given equivalence class?
(global property)



The Sparse Candidate algorithm

Learning a network structure = solving optimization problem in the space of DAGs
⇒ Efficient search algorithms

Sparse Candidate algorithm (Friedman *et. al.*):

- Identify a small number of **candidate parents**,
(simple local statistics like correlation)
- **restrict search** to networks in which only the candidate parents of a variable can be its parents,
- **iteratively adapt** candidate set during search.



Results

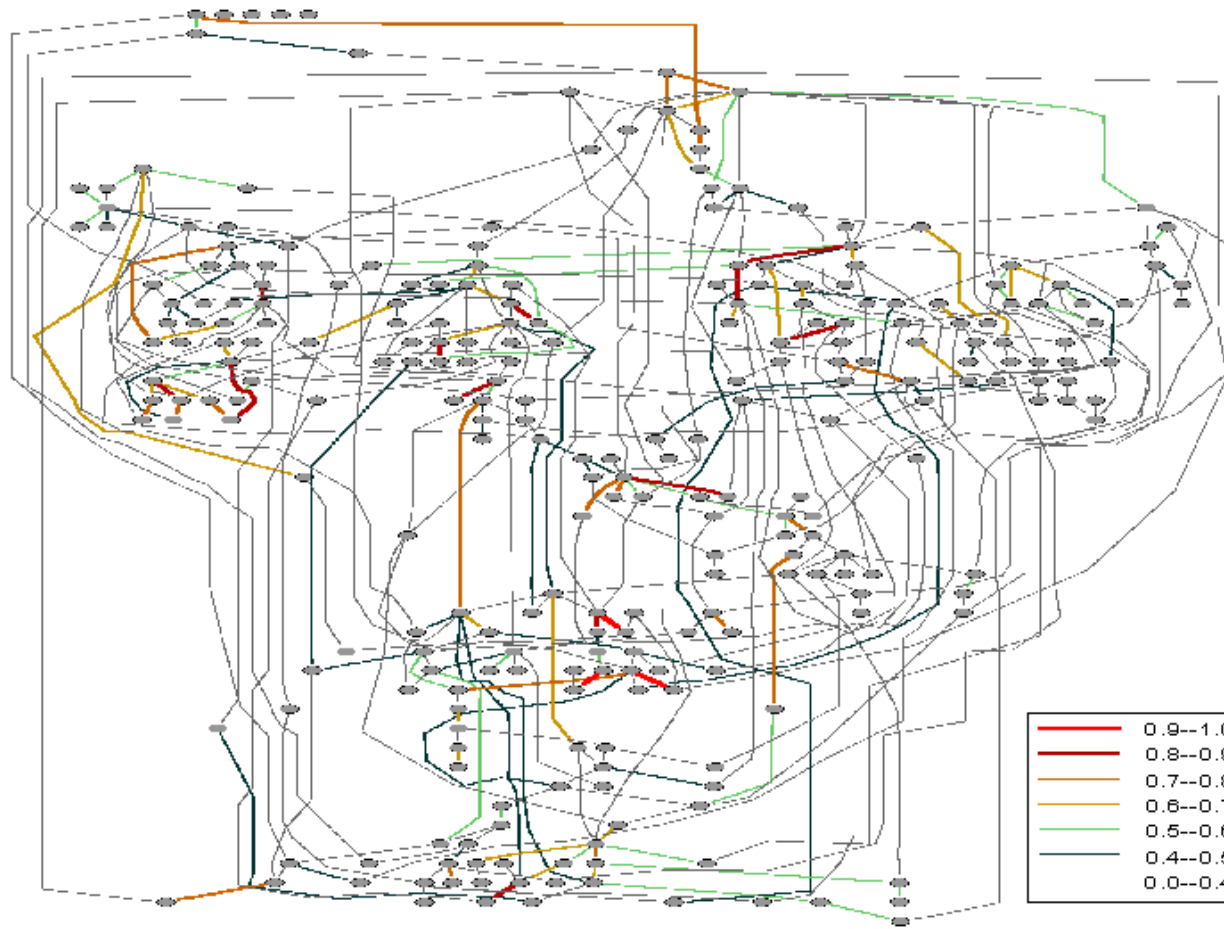
- **Order relations:** only few genes dominate the order (i. e. appear before many genes). These were found to be key genes in the cell-cycle process.
- **Markov relations:** top scoring markov relations between genes were found to indicate a relation in biological function.
- **In general:** Friedman *et. al.* emphasize that BN provide us with a tool that allows biologically plausible conclusions from the data.



Don't get confused!

7

Network Learned



Ideas

- Incorporating biological knowledge
- *Interventional* instead of *observational* data
- How to learn causality from knockout experiments?
- Design of knockout experiments?

