



GILLES KRATZER, APPLIED STATISTICS GROUP, UZH

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# BAYESIAN NETWORKS I LEARNING IN A

# OUTLINE

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- Motivational examples
- Elements of graph theory/probability theory
- Bayesian Network Learning
  - Constraint-based algorithms
  - Score-and-search
- Causal versus acausal thinking
- Real-data applications using R

## Credit Card Fraud Detection Using Bayesian and Neural Networks

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### Abstract

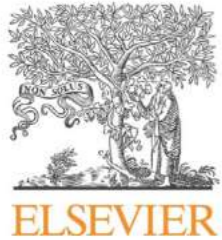
This paper discusses credit card fraud detection by means of digitalization, the great importance of two machine learning under uncertainty

experiment	$\pm 10\%$ false pos	$\pm 15\%$ false pos
ANN-fig 2(a)	60% true pos	70% true pos
ANN-fig 2(a)	47% true pos	58% true pos
ANN-fig 2(c)	60% true pos	70% true pos
BBN-fig 2(e)	68% true pos	74% true pos
BBN-fig 2(g)	68% true pos	74% true pos

Table 1: This table compares the results achieved with ANN and BBN, for a false positive rate of respectively 10% and 15%.

process of learning, able to correctly classify seen before as fraudulent some features of that as follows: first we main of credit card and 4 we briefly ex-

# MOTIVATIONAL EXAMPLE: VETERINARY EPIDEMIOLOGY DATA VISUALISATION



Contents lists available at SciVerse ScienceDirect

## Preventive Veterinary Medicine

journal homepage: [www.elsevier.com/locate/prevetmed](http://www.elsevier.com/locate/prevetmed)



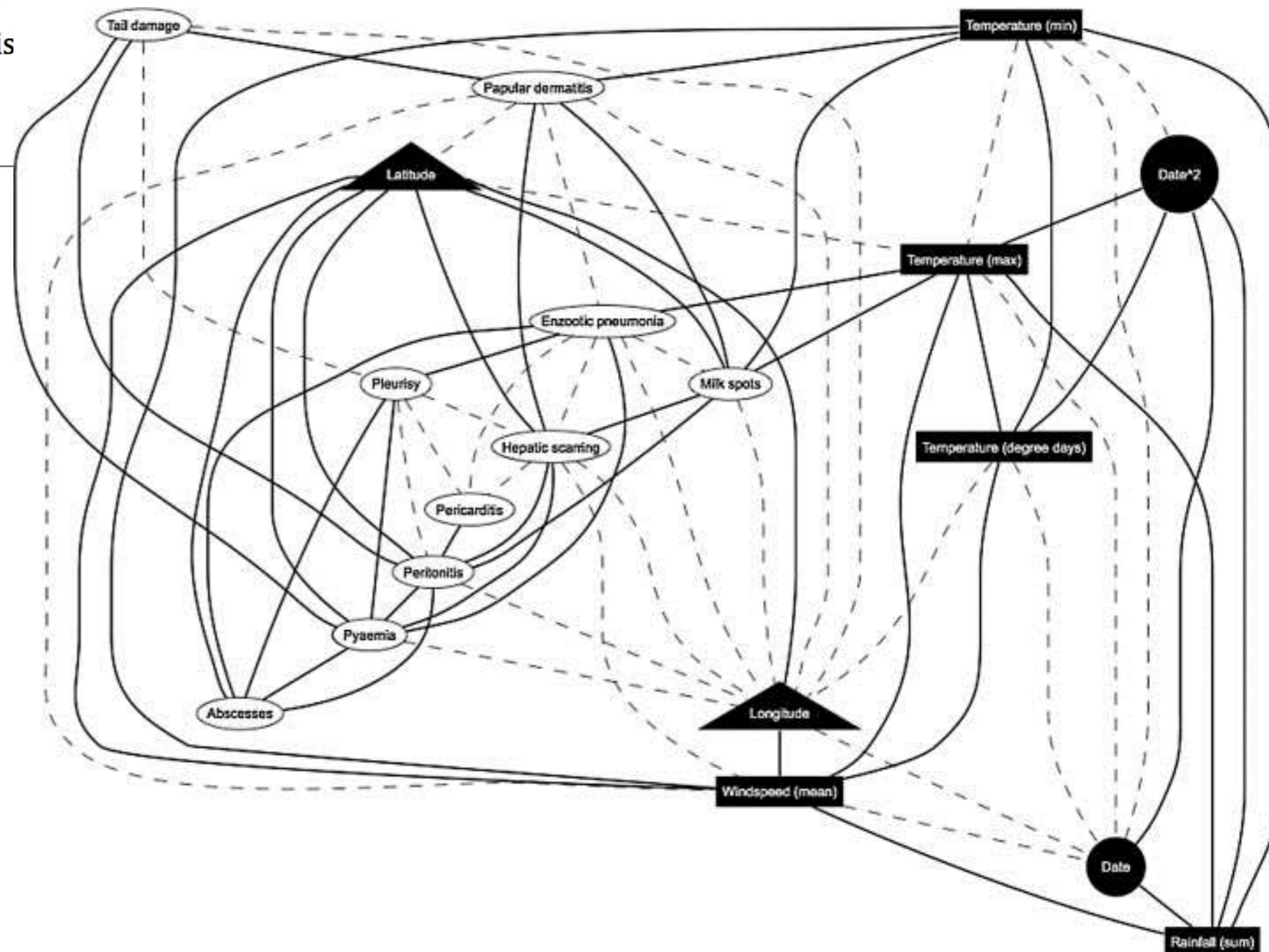
## Using Bayesian networks to explore the role of weather as a potential determinant of disease in pigs

B.J.J. McCormick<sup>a</sup>, M.J. Sanchez-Vazquez<sup>b</sup>, F.I. Lewis

<sup>a</sup> Fogarty International Center, National Institutes of Health, Bethesda, MD 20892, USA

<sup>b</sup> OIE Organisation Mondiale de la Santé Animale, 12, rue de Prony, 75017 Paris, France

<sup>c</sup> Section of Epidemiology, University of Zurich, Zurich, Switzerland



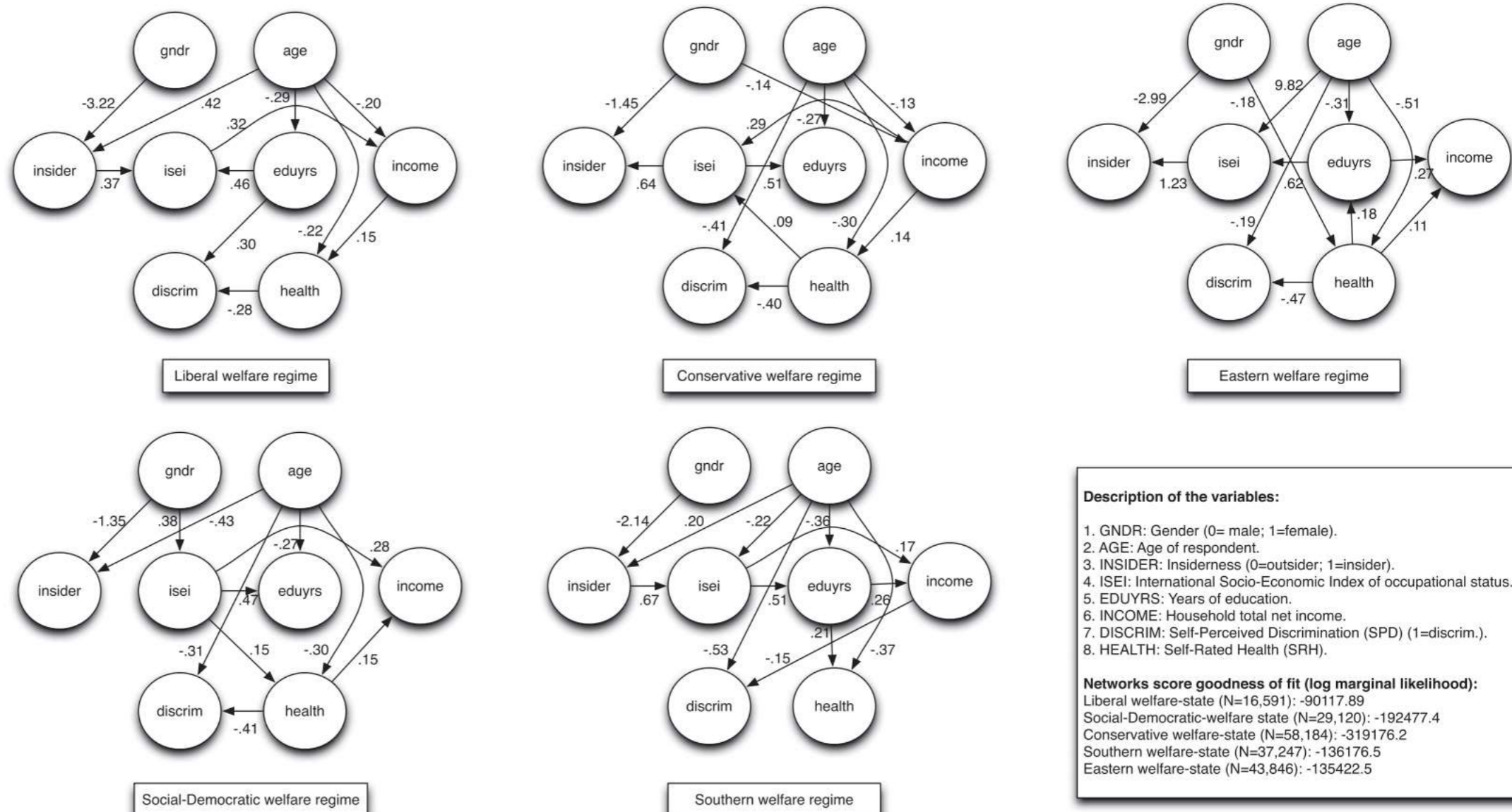
# MOTIVATIONAL EXAMPLE: SOCIAL SCIENCES DATA INTERPRETATION

## Discovering complex interrelationships between socioeconomic status and health in Europe: A case study applying Bayesian Networks

Javier Alvarez-Galvez <sup>a, b, \*</sup>

<sup>a</sup> Loyola University Andalusia, Department of International Studies, Campus de Palmas Altas, Faculty of Political Sciences and Law, Seville 41014, Spain

<sup>b</sup> Complutense University of Madrid, Department of Sociology IV (Research Methodology and Communication Theory), Campus de Somosaguas, Faculty of Political



**Fig. 1.** Bayesian networks describing interrelationships between SES and health in five European welfare states.

# BAYESIAN NETWORKS IN THE MACHINE LEARNING WORLD

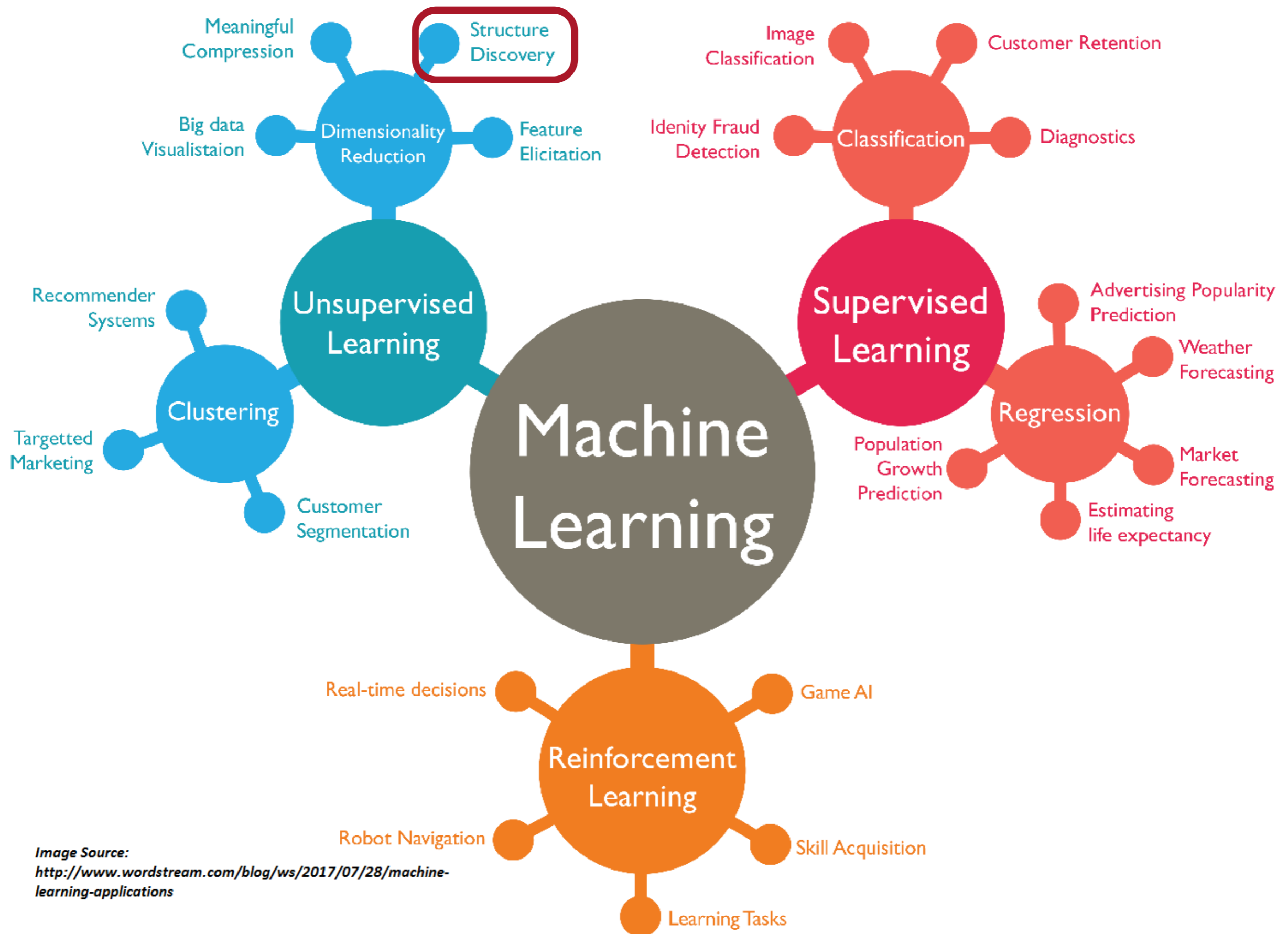


Image Source:  
<http://www.wordstream.com/blog/ws/2017/07/28/machine-learning-applications>

# WHAT IS A BAYESIAN NETWORK?

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Bayesian Networks are defined by two elements:

## Network structure:

Directed Acyclic Graph (**DAG**):  $G = (V, A)$

in which each node  $v_i \in V$  corresponds to a random variable  $X_i$

## Probability distribution:

Probability distribution  $X$  with parameters  $\Theta$ , which can be factorised into smaller local probability distributions according to the arcs  $a_{ij} \in A$  present in the graph.

A BN encodes the factorisation of the joint distribution

$$P(\mathbf{X}) = \prod_{j=1}^n P(X_j \mid \mathbf{Pa}_j, \Theta_j), \text{ where } \mathbf{Pa}_j \text{ is the set of parents of } X_j$$

## SOME ELEMENTS OF PROBABILITY THEORY

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The **conditional probability** of A given B is:  $P(A \mid B) = \frac{P(A, B)}{P(B)}$

**Bayes theorem:**  $P(A \mid B) = \frac{P(B \mid A)P(A)}{P(B)}$

Let A, B and C non intersecting subsets of nodes in a DAG G

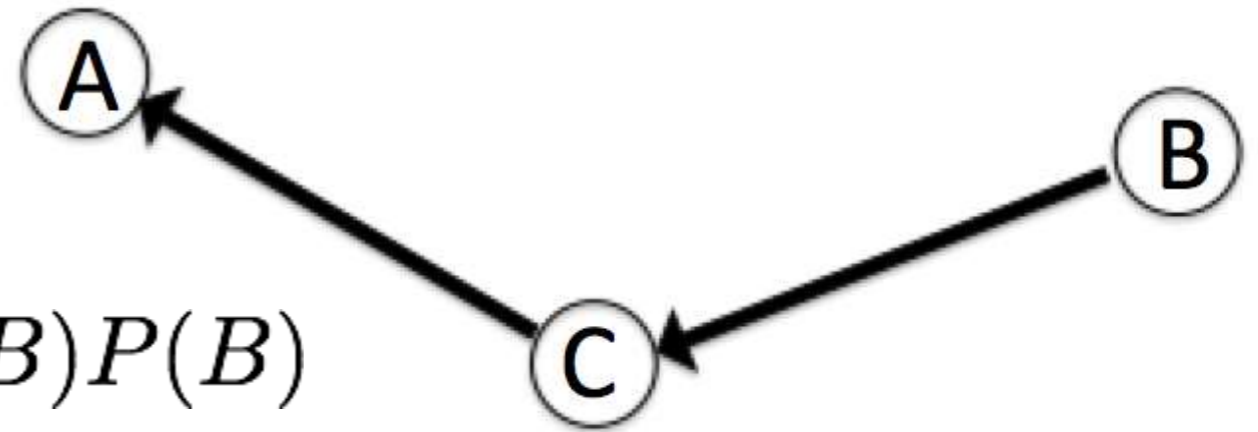
A is **conditionally independent** of B given C if:  $A \perp\!\!\!\perp_P B \mid C$

$$P(A, B \mid C) = P(A \mid C)P(B \mid C)$$

Let A, B and C non intersecting subsets of nodes in a DAG G

A is **conditionally independent** of B given C if:  $A \perp_P B | C$

$$P(A, B | C) = P(A | C)P(B | C)$$



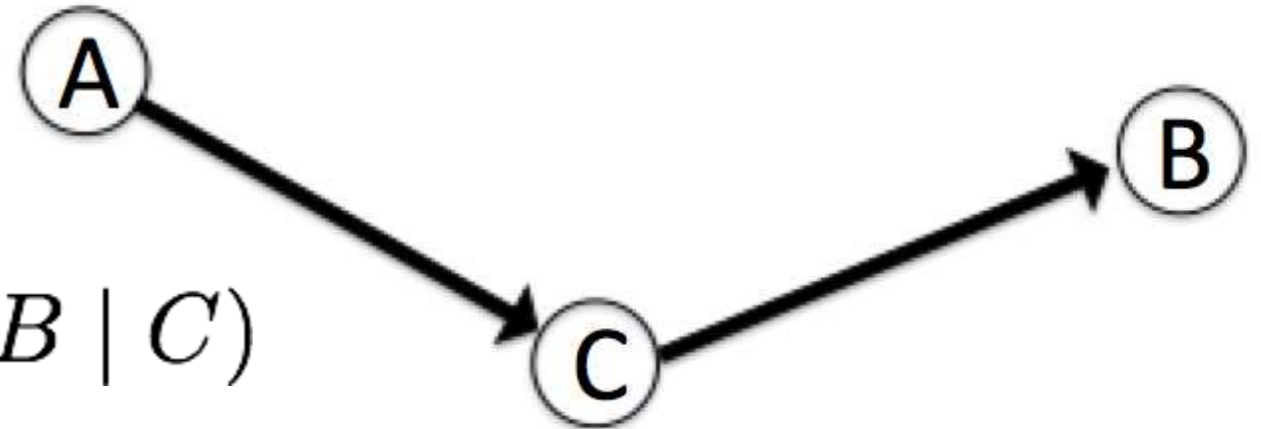
$$P(A, B, C) = P(A | C)P(C | B)P(B)$$

$$\begin{aligned} P(A, B | C) &= \frac{P(A | C)P(C | B)P(B)}{P(C)} \\ &= \frac{P(A | C)P(B, C)}{P(C)} \\ &= P(A | C)P(B | C) \end{aligned}$$

Let A, B and C non intersecting subsets of nodes in a DAG G

A is **conditionally independent** of B given C if:  $A \perp\!\!\!\perp_P B | C$

$$P(A, B | C) = P(A | C)P(B | C)$$



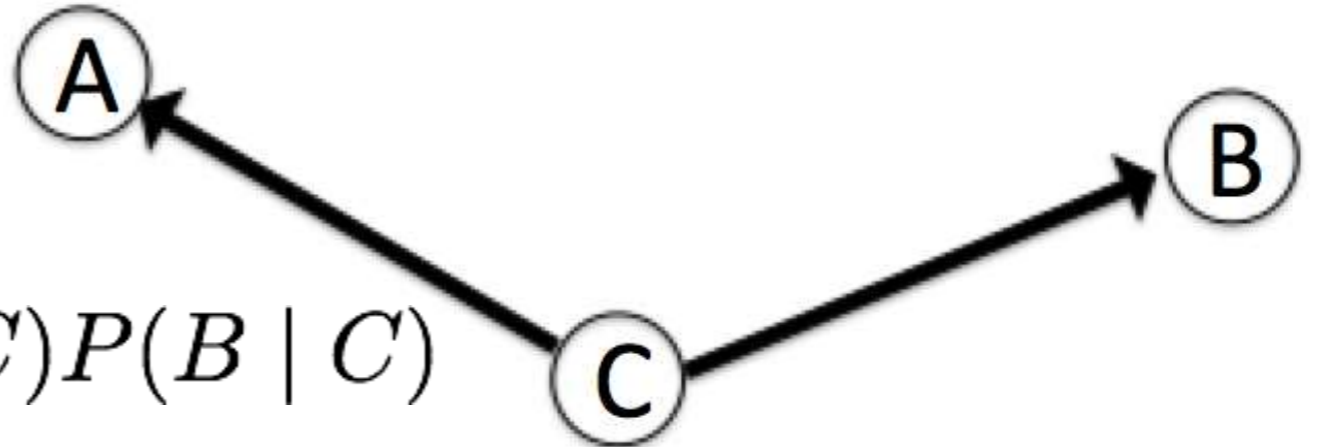
$$P(A, B, C) = P(A)P(C | A)P(B | C)$$

$$\begin{aligned}
 P(A, B | C) &= \frac{P(A)P(C | A)P(B | C)}{P(C)} \\
 &= \frac{P(A, C)P(B | C)}{P(C)} \\
 &= P(A | C)P(B | C)
 \end{aligned}$$

Let A, B and C non intersecting subsets of nodes in a DAG G

A is **conditionally independent** of B given C if:  $A \perp\!\!\!\perp_P B | C$

$$P(A, B | C) = P(A | C)P(B | C)$$



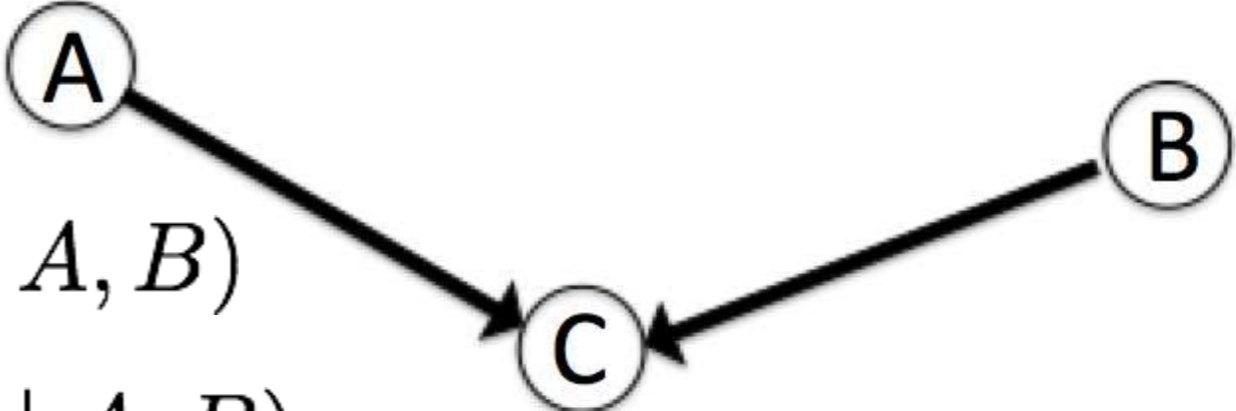
$$P(A, B, C) = P(C)P(A | C)P(B | C)$$

$$\begin{aligned} P(A, B | C) &= \frac{P(C)P(A | C)P(B | C)}{P(C)} \\ &= P(A | C)P(B | C) \end{aligned}$$

Let  $A$ ,  $B$  and  $C$  non intersecting subsets of nodes in a DAG  $G$

$A$  is **conditionally independent** of  $B$  given  $C$  if:  $A \perp_P B | C$

$$P(A, B | C) = P(A | C)P(B | C)$$


$$\begin{aligned} P(A, B, C) &= P(A)P(B)P(C | A, B) \\ P(A, B | C) &= \frac{P(A)P(B)P(C | A, B)}{P(C)} \\ &= \frac{P(A)P(B)P(A, B, C)}{P(A)P(B)P(C)} \\ &= P(A, B | C) \end{aligned}$$

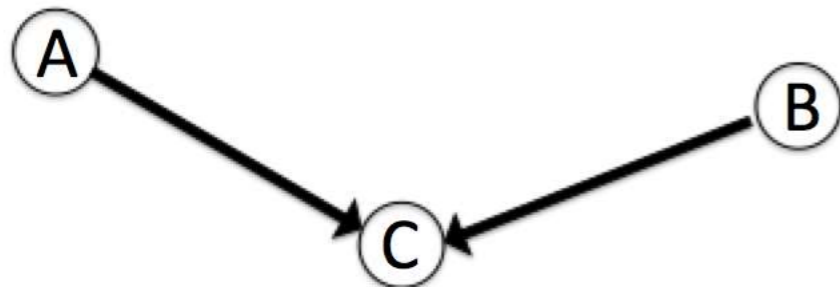
$A \not\perp_P B | C$

Let A, B and C non intersecting subsets of nodes in a DAG G

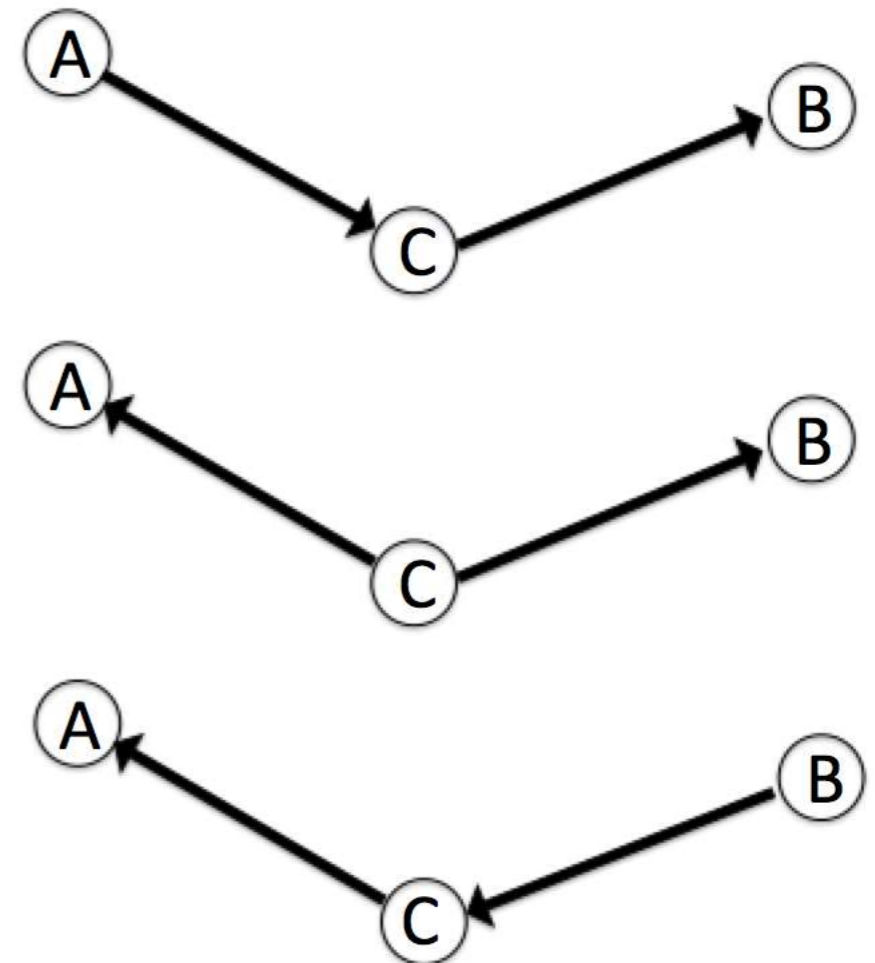
A is **conditionally independent** of B given C if:  $A \perp_P B | C$

$$P(A, B | C) = P(A | C)P(B | C)$$

$$A \not\perp_P B | C$$

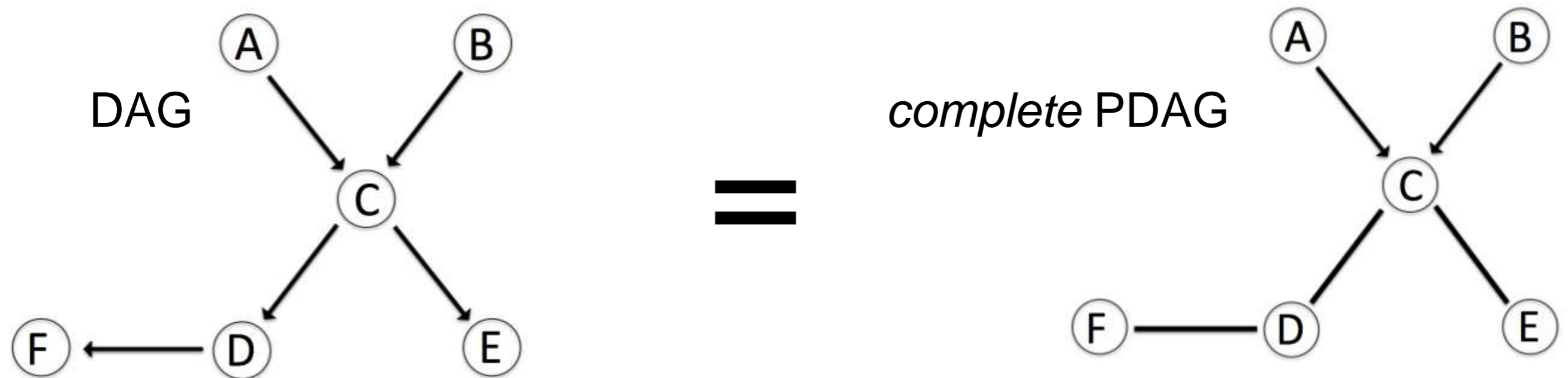


$$A \perp_P B | C$$



# LEARNING BAYESIAN NETWORKS

- ▶ In a practical perspective, for **observational** data, if learning algorithms rely on **probabilistic learning algorithm**. Then one can learn up to the **Markov equivalence class**.
- ▶ **Markov equivalence class** are the set of DAGs that have the same **skeleton** and **v-structure**.



- ▶ the arrows on the path meet either head-to-tail or tail-to-tail at the node, and the node is in the set C, or
- ▶ the arrows meet head-to-head at the node, and neither the node, nor any of its descendants, are C.

If all paths from A to B are blocked, A is said to be **d-separated** from B by C.

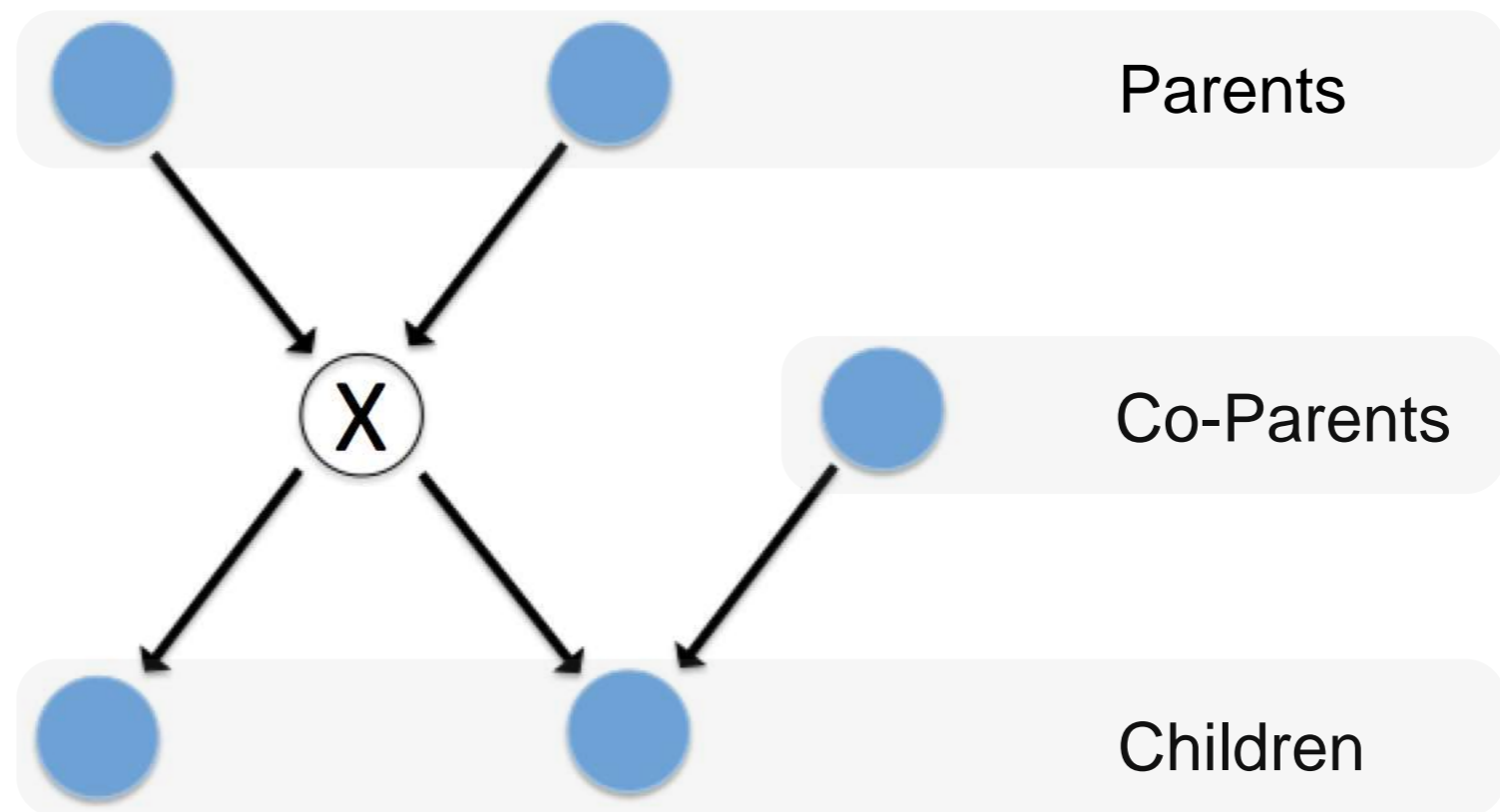
**Theorem** ([Verma & Pearl, 1988](#)): A is d-separated from B by C if, and only if, the

$$A \perp\!\!\!\perp_G B | C$$

joint distribution over all variables in the graph satisfies:

# ELEMENT OF GRAPH THEORY

The **Markov Blanket** of a node is the set of **parents**, **co-parents** and **children**.



$$P(X_k \mid X_n, k \neq n) = P(X_k \mid X_{\text{MB}(k)}), \forall k$$

The **Markov Blanket** of a node is the set of nodes that **shields** the index node from the res

$$\mathcal{M} = (\mathcal{S}, \theta_{\mathcal{M}})$$

Model selection

Structure learning

Parameter estimation

Parameter learning

$$P(\mathcal{M}|\mathcal{D}) = \underbrace{P(\theta_{\mathcal{M}}, \mathcal{S}|\mathcal{D})}_{\text{model learning}} = \underbrace{P(\theta_{\mathcal{M}}|\mathcal{S}, \mathcal{D})}_{\text{parameter learning}} \underbrace{P(\mathcal{S}|\mathcal{D})}_{\text{structure learning}}$$

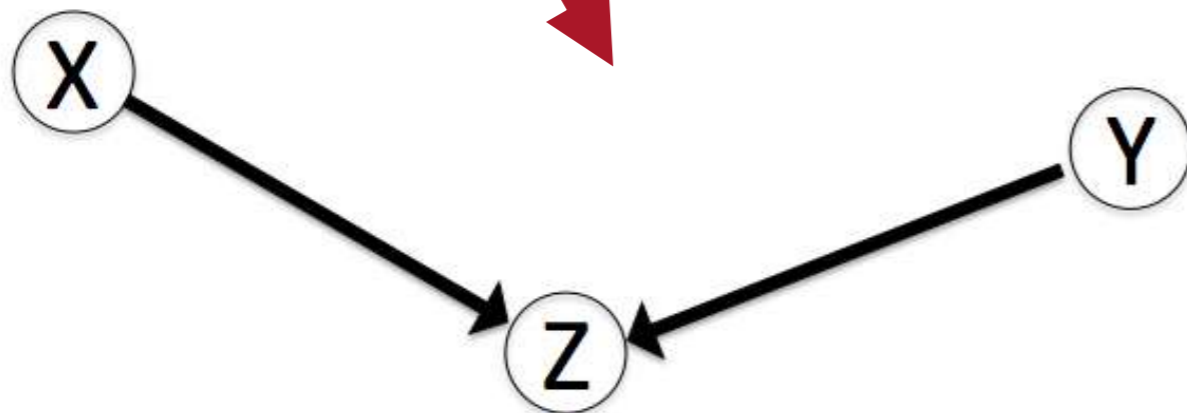
	Fully Observed data	Missing data/hidden variables
Known graph structure	Easy Sample statistics	EM algorithm Gradient ascent Variational inference Doable
Unknown graph structure	Doable Search-and-score PC algorithm	Hard Structural EM

## Constraint based algorithms

$$P_{X \perp\!\!\!\perp Y|Z} < \alpha$$



$$X \perp\!\!\!\perp_s Y|Z = X \perp\!\!\!\perp Y|Z$$



## Search-and-score algorithms

Maximum a posteriori score

$$G^* = \operatorname{argmax}_G f(\mathcal{D}, G, n, \dots)$$

Example of scoring functions:

- Bayesian versus ML scores
  - log marginal likelihood
  - Bayesian-Dirichlet (BDeu, BDs, BDe)
  - Bayesian Information Criterion (BIC)

## Constraint-based algorithms

- ▶ *Inductive Causation (IC)*: ([Verma and Pearl, 1991](#))
  - ▶ Provides a framework for learning the structure of Bayesian networks using conditional independence tests in three steps
  - ▶ A major problem of the IC algorithm is that the first two steps cannot be applied to any real-world problem due to computational complexity ...
- ▶ *PC*: first practical application of the IC algorithm ([Spirtes et al., 2001](#))
  - ▶ backward selection procedure from the saturated graph
- ▶ *Grow-Shrink (GS)* ([Margaritis, 2003](#))
  - ▶ Simple forward selection MB detection approach
- ▶ *Incremental Association (IAMB)*: ([Tsamardinos et al., 2003](#))
  - ▶ two-phase selection scheme based on a forward selection followed by a backward selection of the MB

# LEARNING BAYESIAN NETWORKS REQUIRES A Markov and faithfulness assumption

- Conditional independencies in the distribution exactly equal the ones encoded in the DAG via **d-separation**

$$A \perp\!\!\!\perp_G B|C \begin{array}{c} \text{Markov} \\ \xleftrightarrow{\hspace{1cm}} \\ \text{Faithful} \end{array} A \perp\!\!\!\perp_P B|C$$

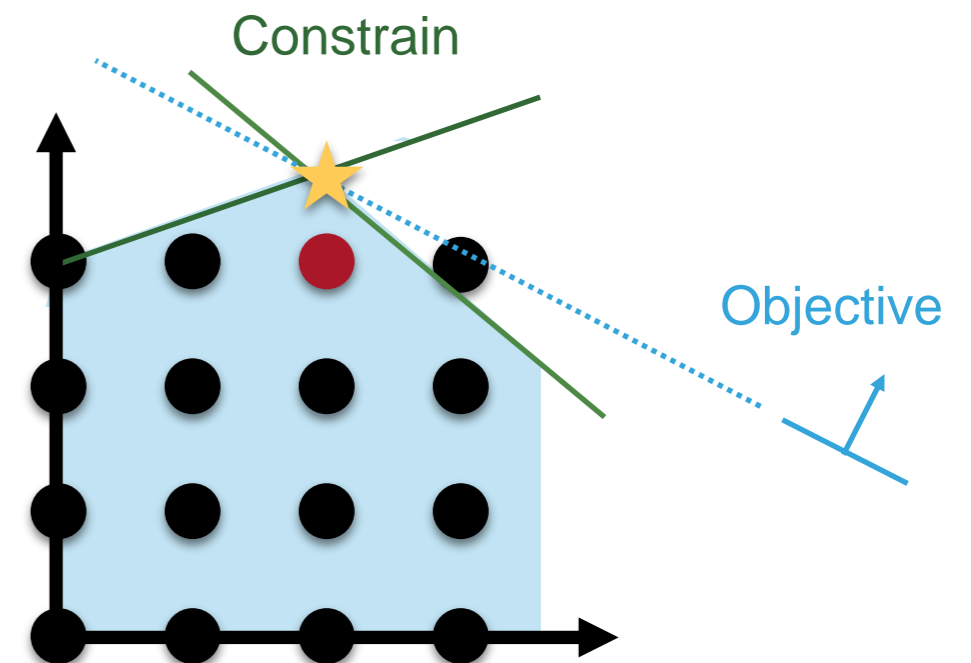
- Causal sufficiency**: no unmeasured common causes

In a practical perspective:

- Testing mixture of data?
- Testing assumptions?

## Score-and-search algorithms

- *Heuristic approaches / Greedy search*
  - Hill-climbing (with possibly random restarts/stochastics ... )
  - Tabu search ([Glover, 1986](#))
  - Simulated annealing ([Kirkpatrick et al, 1983](#))
  - Plus an entire zoo of methods ...
- *Exact search*
  - Exact node ordering ([Koivisto et al. , 2004](#))
  - Learning with cutting planes ([Cussens, 2012](#))



# LEARNING BAYESIAN NETWORKS

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## Scores

- Decomposability!
- Discrete BNs:
  - Bayesian-Dirichlet: **BDeu** ([Heckerman et al. ,1995](#))
- Score equivalence for additive regression framework:
  - **Bayesian based scores**: not always score equivalent due to the prior!
  - **Information theoretic scores**: BIC asymptotically score equivalent

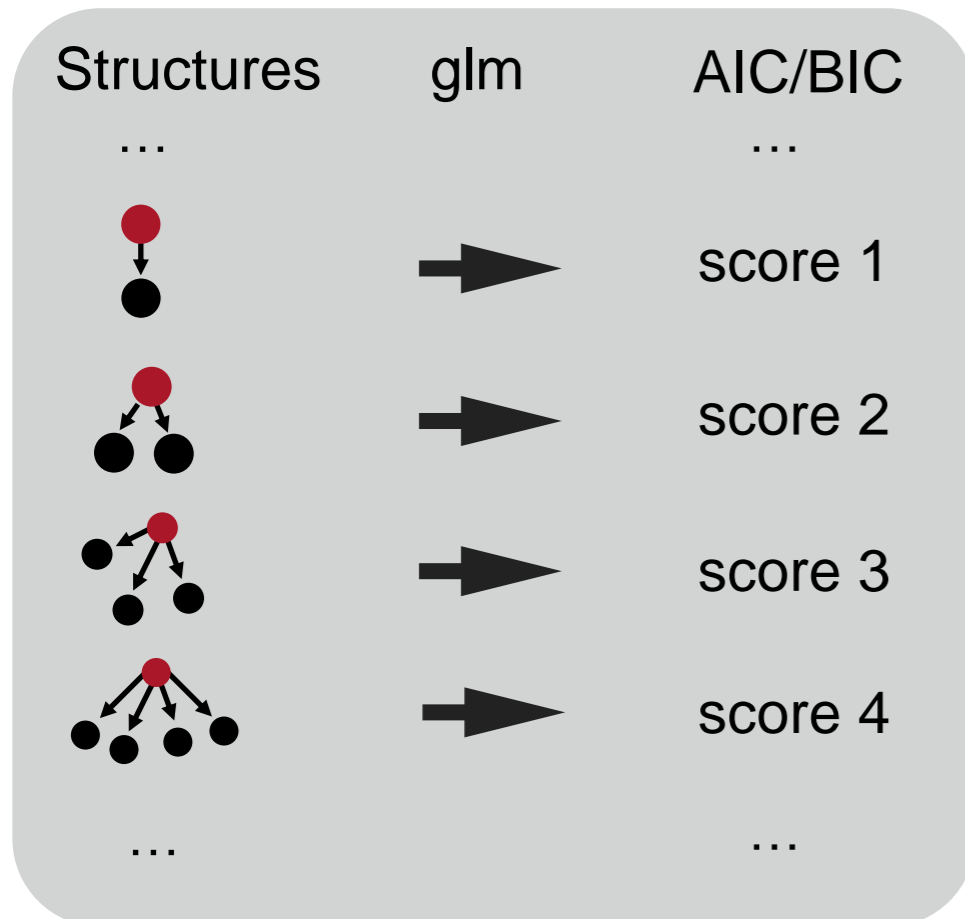
## Counter example

- Maximum likelihood estimator ... return fully connected BN!

## In a practical perspective:

- Scoring mixture of data?
- Score equivalence!

## Search and score algorithm

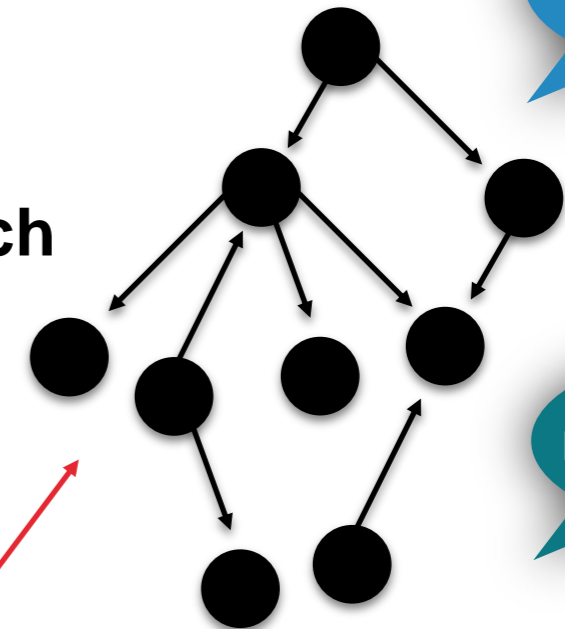


Exact or heuristic search



**Causality!**

*Ban/Retain  
structures*



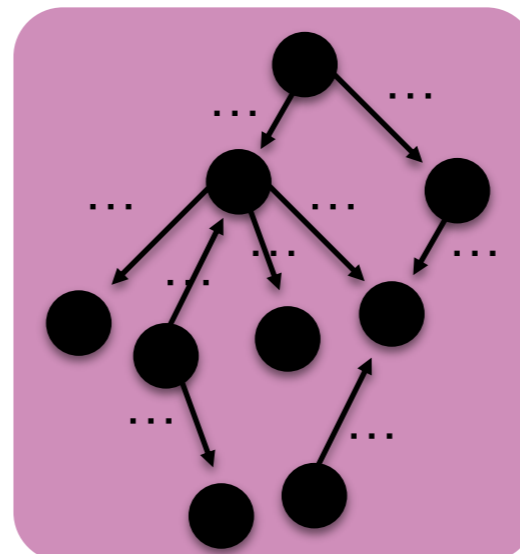
Adjustment

Random effect

Bayesian network with  
highest posterior  
probability

## Parameter estimation

- › compute marginal posterior density
- › regression estimate



**Using R**

`buildscorecache()`

`mostprobable()`

`fitabn()`

# CAUSAL THINKING VERSUS ACAUSAL THINKING

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- Strong assumptions ... but common in statistics, no?
- *“It seems that if conditional independence judgements are byproducts of stored causal relationships, then tapping and representing those relationships directly would be a more natural and more reliable way of expressing what we know or believe about the world. This is indeed the philosophy behind causal Bayesian networks.”* (Pearl, 2009)
- The **do-calculus**
  - **Interventions**
  - In epidemiology: **Randomised Controlled Trial**
- So ... BN is a nice framework to treat causal and causal thinking

# R CODE: SOFTWARE IMPLEMENTATION

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Popular R packages (available on [CRAN](#))

## **bnlearn**

- Learning via constraint-based and score-based algorithms (many!)

## **pcalg**

- Robust estimation of CPDAG via the PC-Algorithm

## **deal**

- Learning BNs with mixed (discrete and continuous) variables

## **catnet**

- Discrete BNs using likelihood-based criteria

## **abn**

- Learning BNs with mixed (discrete, continuous, Poisson) variables
- Score based methods: Bayesian and frequentist estimation
- Exact and heuristic search

**Disclaimer:** I am author and maintainer of the abn R package. I will use it for the example part.

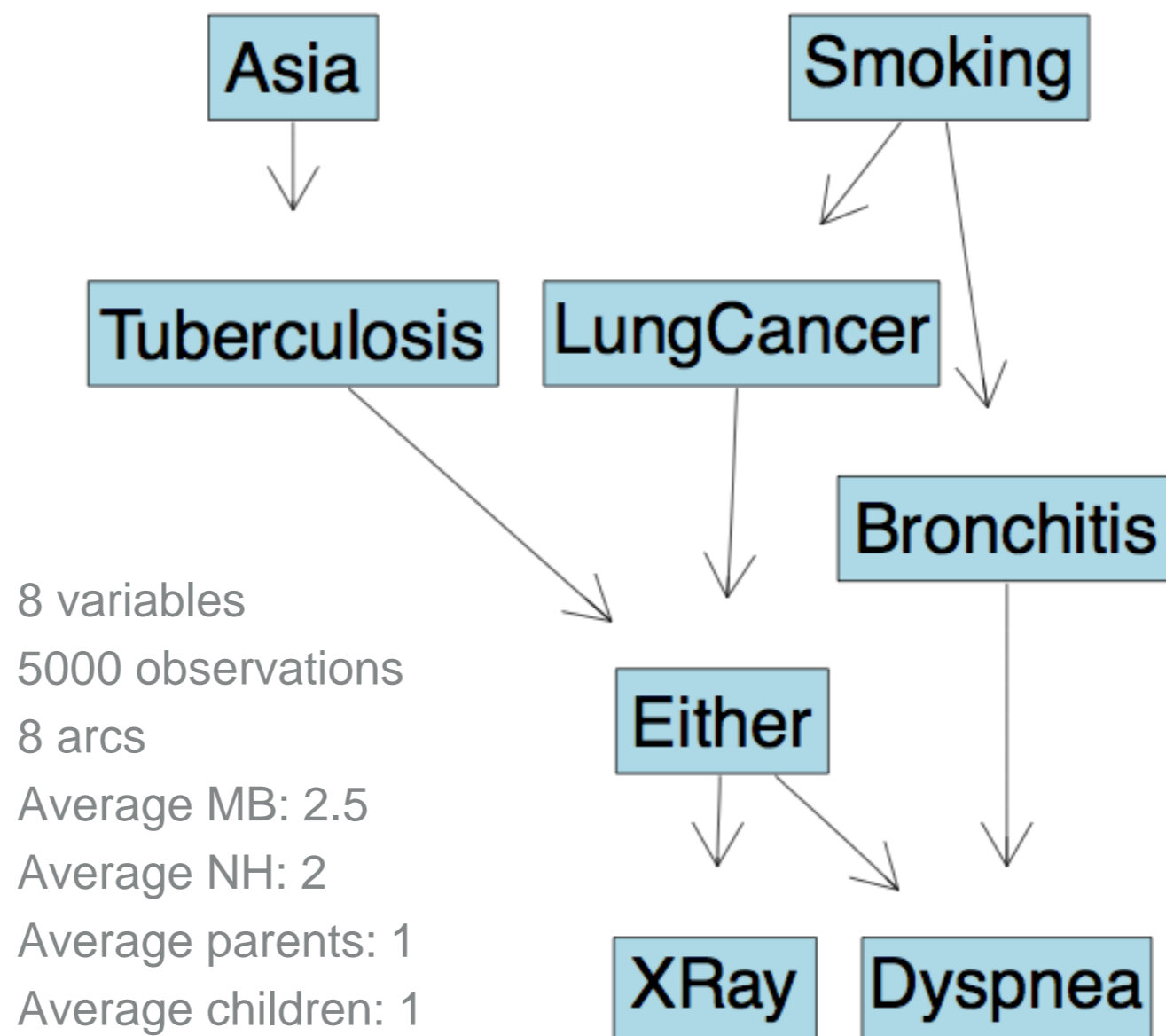
# R CODE: EXAMPLE ASIA

Proposed by [Lauritzen et al., 1988](#) and provided by [Scutari, 2009](#)

“Shortness-of-breath (*dyspnoea*) may be due to *tuberculosis*, *lung cancer* or *bronchitis*, or none of them, or more than one of them. A recent visit to *Asia* increases the chances of *tuberculosis*, while *smoking* is known to be a risk factor for both *lung cancer* and *bronchitis*. The results of a single chest *X-ray* do not discriminate between *lung cancer* and *tuberculosis*, as neither does the presence or absence of *dyspnoea*.”

```
##defining distributions
dist = list(Asia = "binomial",
            Smoking = "binomial",
            Tuberculosis = "binomial",
            LungCancer = "binomial",
            Bronchitis = "binomial",
            Either = "binomial",
            XRay = "binomial",
            Dyspnea = "binomial")

#plot BN
plotabn(dag.m = ~Asia|Tuberculosis +
        Tuberculosis|Either +
        Either|XRay:Dyspnea +
        Smoking|Bronchitis:LungCancer +
        LungCancer|Either +
        Bronchitis|Dyspnea,
        data.dists = dist,
        edgedir = "cp",
        fontsize.node = 30,
        edge.arrowwise = 3)
```



# ASIA: SCORE BASED ALGORITHM

```
##=====
##score based algorithm
##=====

#loglikelihood score
bsc.compute <- buildscorecache(data.df = asia,
                               data.dists = dist,
                               max.parents = 2)

dag <- mostprobable(score.cache = bsc.compute)
plotabn(dag.m = dag, data.dists = dist, fontsize.node = 30, edge.arrows = TRUE)
```

```
> compareDag(ref = t(dag.adj),
+            test = dag)

$TPR
[1] 0.75

$FPR
[1] 0.01785714

$Accuracy
[1] 0.953125

$FDR
[1] 0.2857143

$`G-measure`
[1] 0.8017837

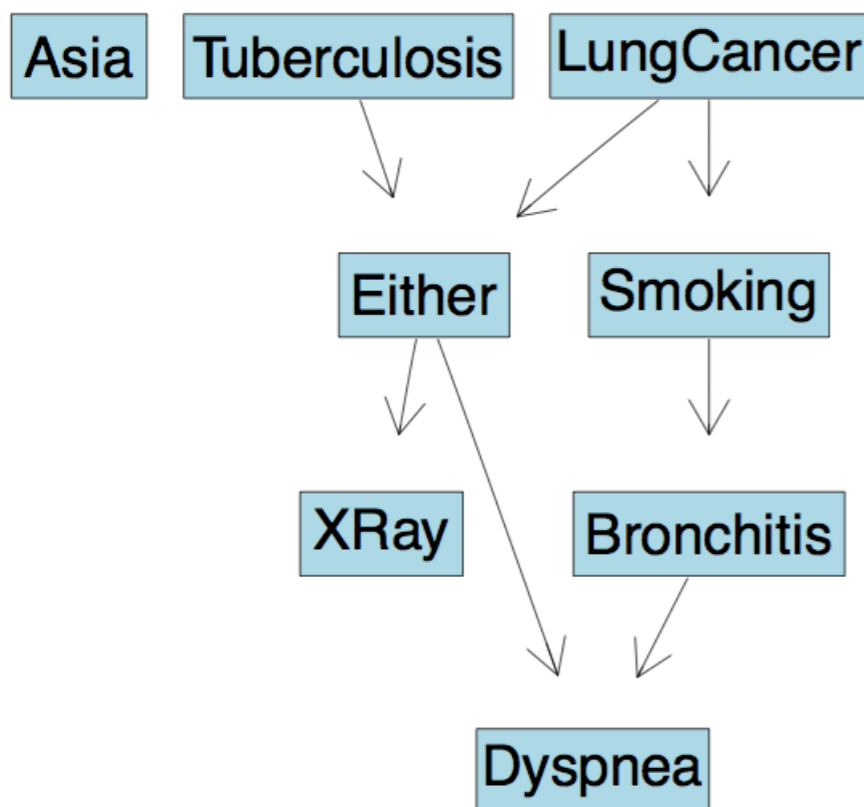
$`F1-score`
[1] 44.8

$PPV
[1] 0.8571429

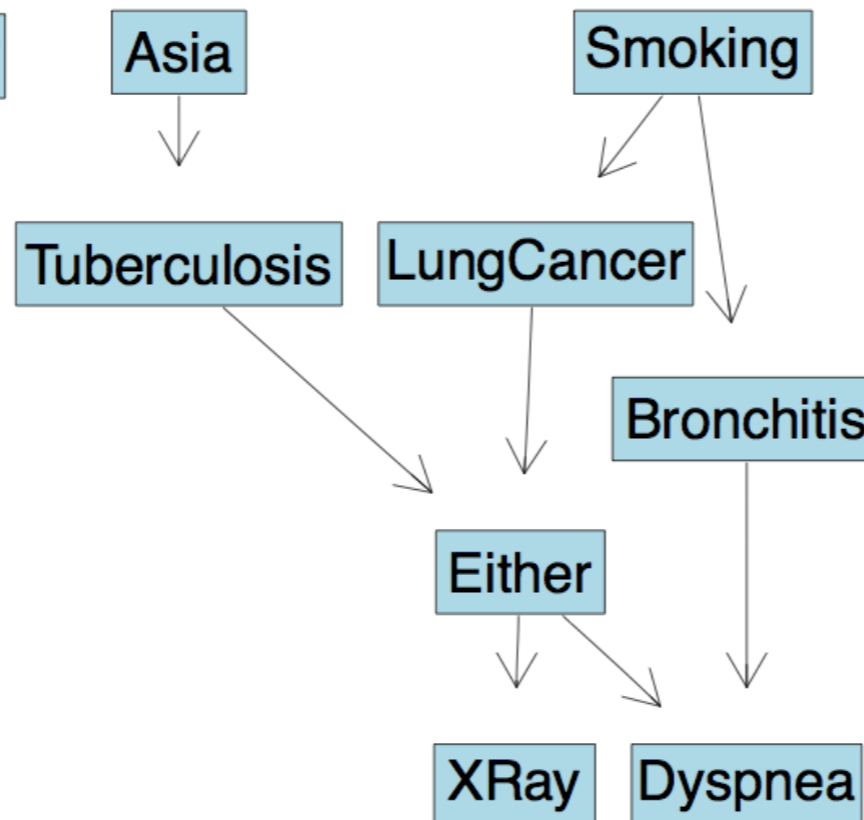
$FOR
[1] 0.2857143

$`Hamming-distance`
[1] 3
```

Learned



Truth



# ASIA: KNOWN NETWORK

```
fitabn(dag.m = ~Asia|Tuberculosis+
  Tuberculosis|Either +
  Either|XRay:Dyspnea +
  Smoking|Bronchitis:LungCancer +
  LungCancer|Either +
  Bronchitis|Dyspnea,data.df = asia,data.dists = dist)$modes
```

```
fitabn.mle(dag.m = dag.adj,data.df = asia,data.dists = dist)$coef
```

```
$Asia
Asia|(Intercept) Asia|Tuberculosis
      -4.811200      1.765763

$Smoking
Smoking|(Intercept) Smoking|LungCancer Smoking|Bronchitis
      -1.027065      2.356988      1.807460

$Tuberculosis
Tuberculosis|(Intercept) Tuberculosis|Either
      -12.22120      10.21823

$LungCancer
LungCancer|(Intercept) LungCancer|Either
      -12.07565      14.18547

$Bronchitis
Bronchitis|(Intercept) Bronchitis|Dyspnea
      -1.388644      3.200393

$Either
Either|(Intercept) Either|XRay Either|Dyspnea
      -8.656348      8.259773      1.538789

$XRay
XRay|(Intercept)
      -2.052496

$Dyspnea
Dyspnea|(Intercept)
      -0.1201444
```

```
$Asia
      Asia|intercept Tuberculosis
[1,]      -4.811371      1.766849

$Smoking
      Smoking|intercept LungCancer Bronchitis
[1,]      -1.027075      2.357079      1.807472

$Tuberculosis
      Tuberculosis|intercept Either
[1,]      -8.517393      6.516139

$LungCancer
      LungCancer|intercept Either
[1,]      -8.517393      10.62598

$Bronchitis
      Bronchitis|intercept Dyspnea
[1,]      -1.388655      3.200415

$Either
      Either|intercept XRay Dyspnea
[1,]      -8.665128      8.268402      1.539146

$XRay
      XRay|intercept
[1,]      -2.0525

$Dyspnea
      Dyspnea|intercept
[1,]      -0.1201443
```

# ASIA: EXTERNAL KNOWLEDGE

```
##=====
##external knowledge
##=====

##recent visit to Asia increases risk of tuberculosis
bsc.compute <- buildscorecache.mle(data.df = asia,
                                   data.dists = dist,
                                   max.parents = 2,
                                   dag.retained = ~Tuberculosis|Asia)

dag <- mostprobable(score.cache = bsc.compute, score = "bic")
plotabn(dag.m = dag, data.dists = dist, fontsize.node = 30, edge.arrow
```

```
> compareDag(ref = t(dag.adj),
+            test = (dag))
$TPR
[1] 0.875

$FPR
[1] 0.01785714

$Accuracy
[1] 0.96875

$FDR
[1] 0.125

$`G-measure`
[1] 0.875

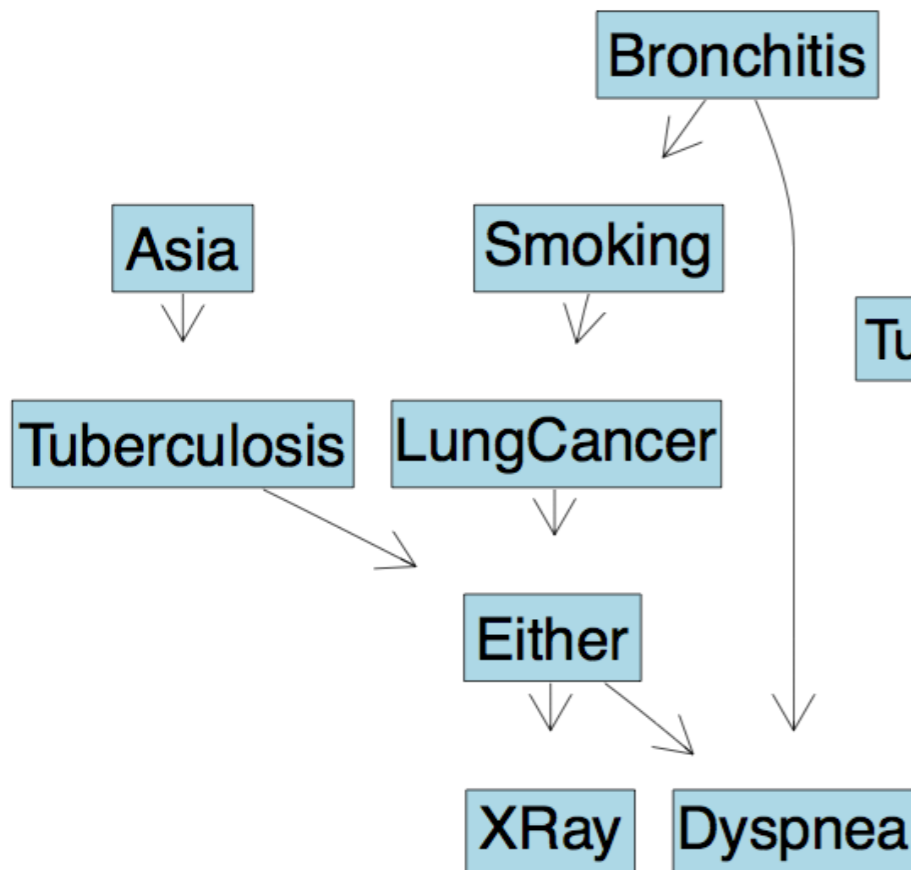
$`F1-score`
[1] 56

$PPV
[1] 0.875

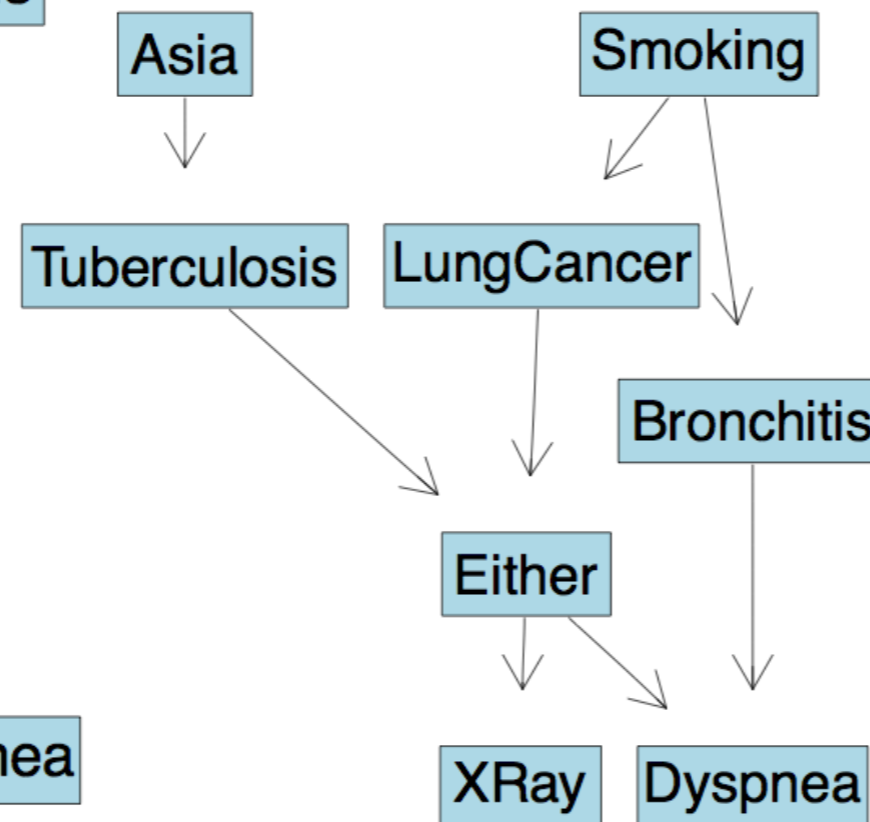
$FOR
[1] 0.125

$`Hamming-distance`
[1] 2
```

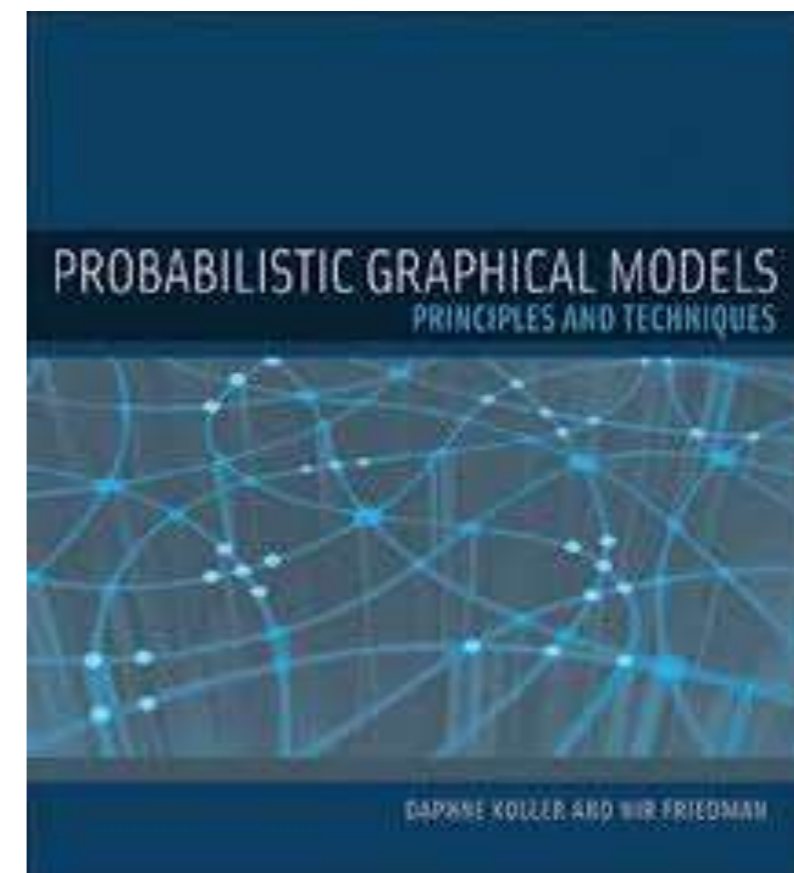
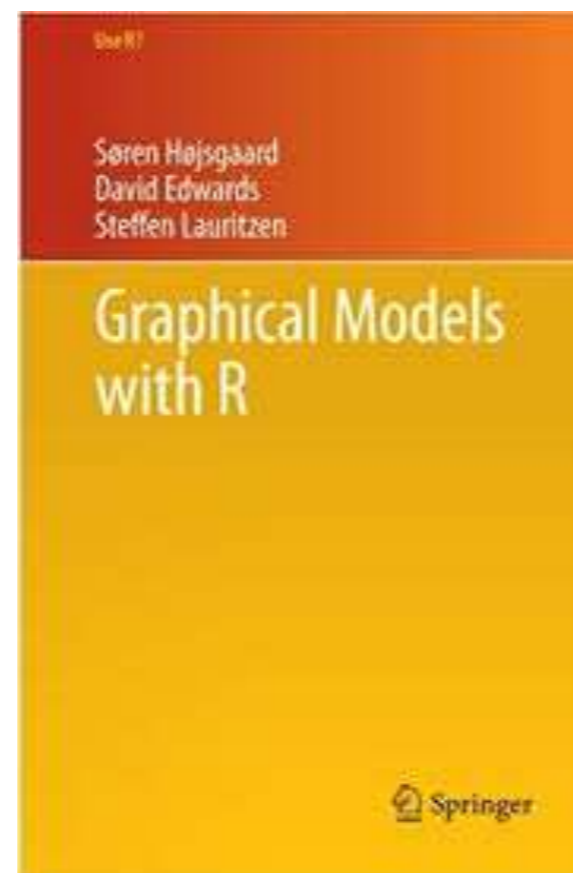
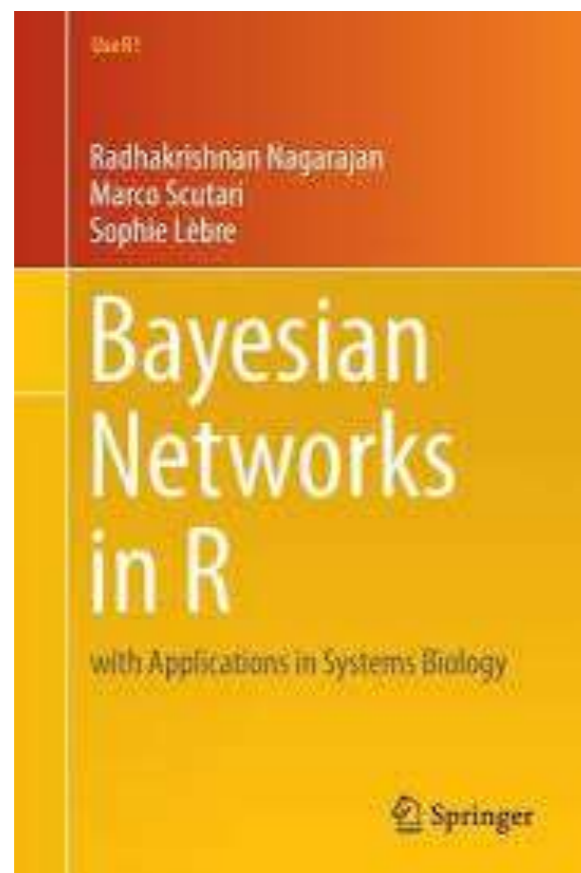
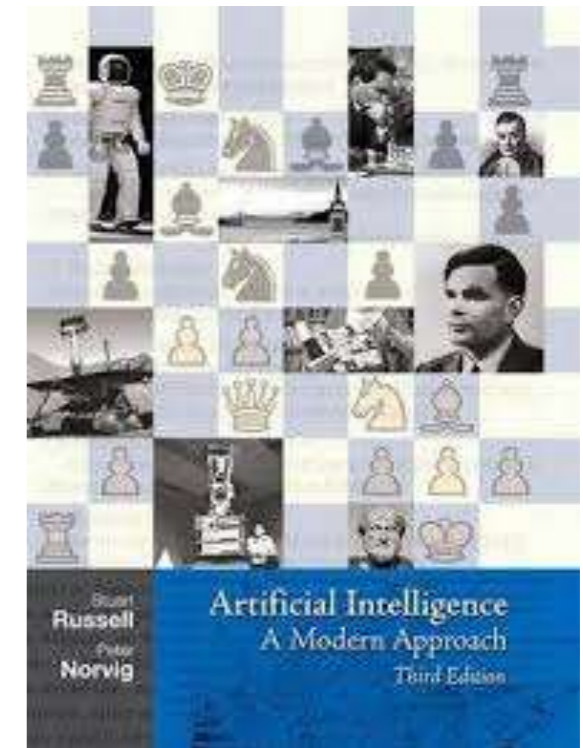
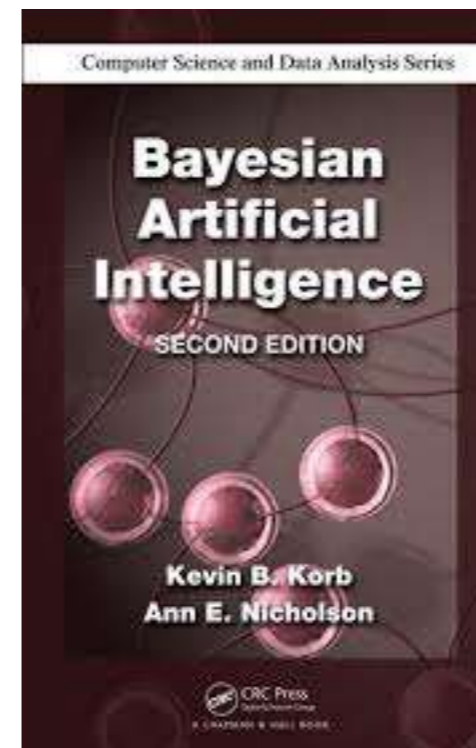
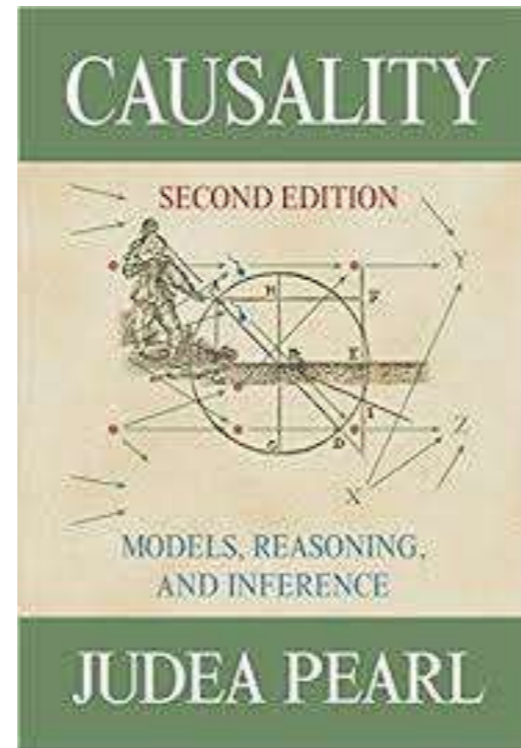
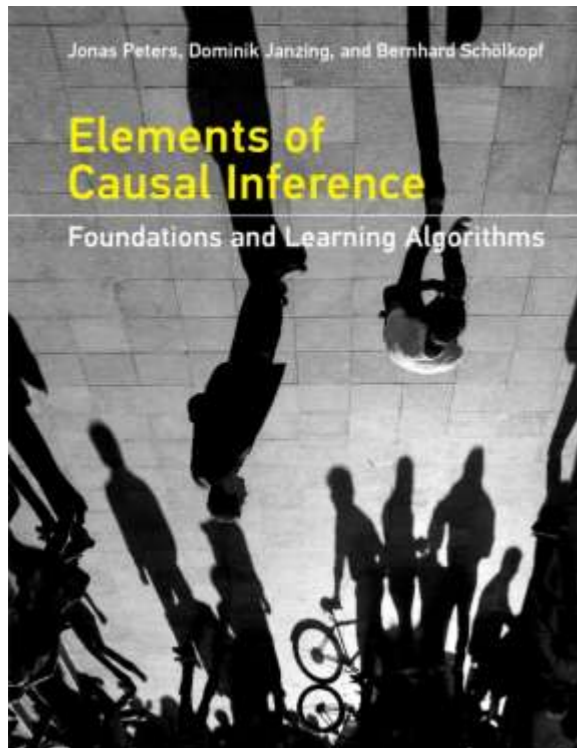
Learned



Truth



# SELECTED BIBLIOGRAPHY



# Thank you for your attention



xkcd.com

## **Backup slides**

# ASIA: BOOTSTRAPPING

```
library(doParallel)
library(foreach)

cl <- makeCluster(2)
registerDoParallel(cl)

set.seed(1120)
nboot <- 200
nvars <- dim(asia)[2]
nobs <- dim(asia)[1]
bootstrap.dag <- array(data = NA, dim = c(nvars, nvars, nboot))

start_time <- Sys.time()
bootstrap.dag <- foreach(i = 1:nboot, .packages = c("mlabn", "abn")) %dopar% {
  mycache.computed.mle <- buildscorecache.mle(data.df = asia[sample(x = 1:nobs, size = 0.8*nobs, replace = FALSE), ],
                                              data.dists = dist,
                                              max.parents = 2,
                                              dry.run = FALSE,
                                              maxit = 1000,
                                              tol = 1e-11)

  dag <- mostprobable(score.cache = mycache.computed.mle, score = "bic")
}
compute_time <- Sys.time()-start_time

##analysis
df.boot <- array(data = unlist(bootstrap.dag), dim = c(8, 8, 200))

dag<-apply(df.boot, 1:2, mean)

#dag.mdl<-dag.before

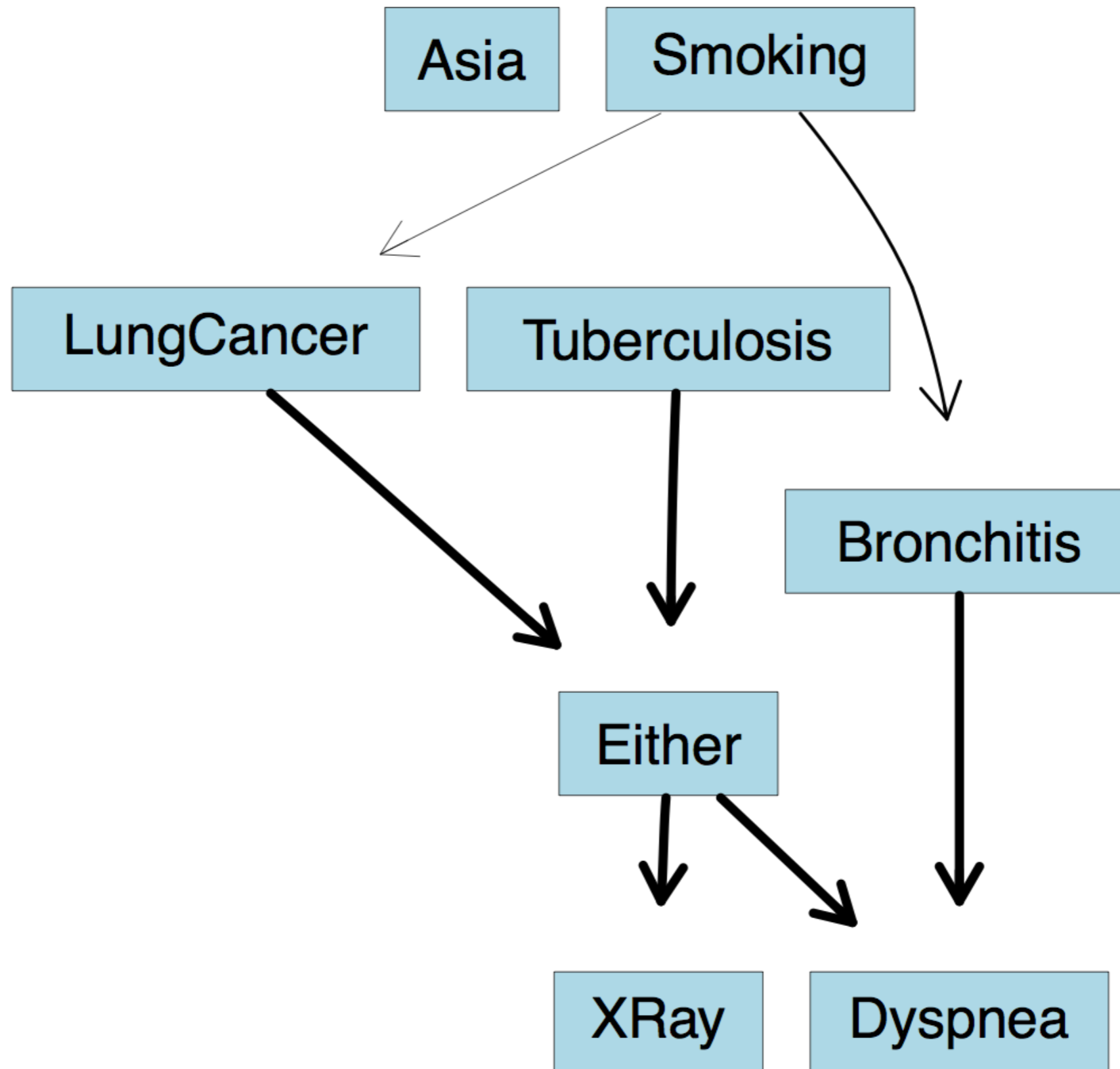
colnames(dag) <- rownames(dag) <- names(dist)

dag.boot.50 <- dag
dag.boot.50[dag>0.5] <- 1
dag.boot.50[dag<=0.5] <- 0

dag[dag<=0.5] <- 0

colnames(dag.boot.50) <- rownames(dag.boot.50) <- names(dist)

plotabn(dag.m = t(dag.boot.50), data.dists = dist, fontsize.node = 30, arc.strength = 10*dag, digit.precision = 2, edge.arrowwise = 3)
```



# ASIA: HOW MANY PARENT ARE NEEDED?

```
res.mlik <- NULL
res.aic <- NULL
res.bic <- NULL
res.mdl <- NULL

for(i in 1:4){
  mycache.computed.mle <- buildscorecache.mle(data.df = asia,
                                              data.dists = dist,
                                              max.parents = i,
                                              dry.run = FALSE,
                                              maxit = 1000,
                                              tol = 1e-11)

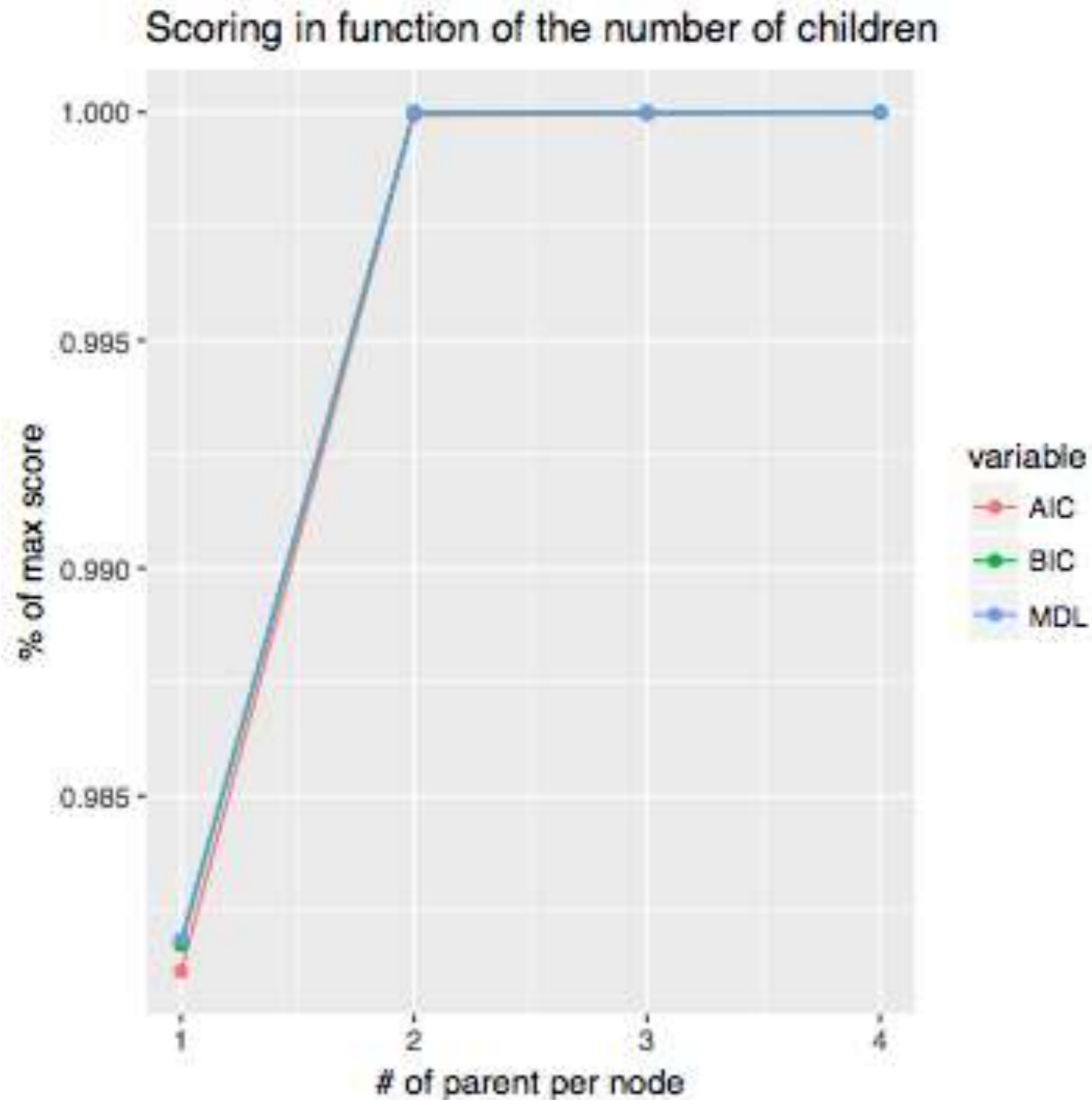
  dag <- mostprobable(score.cache = mycache.computed.mle, score = "aic")
  res.aic <- rbind(res.aic, fitabn.mle(dag.m = dag, data.df = mycache.computed.mle$data.df, data.dists = dist)$aic)
  dag <- mostprobable(score.cache = mycache.computed.mle, score = "bic")
  res.bic <- rbind(res.bic, fitabn.mle(dag.m = dag, data.df = mycache.computed.mle$data.df, data.dists = dist)$bic)
  dag <- mostprobable(score.cache = mycache.computed.mle, score = "mdl")
  res.mdl <- rbind(res.mdl, fitabn.mle(dag.m = dag, data.df = mycache.computed.mle$data.df, data.dists = dist)$mdl)
}

library(ggplot2)
library(reshape)
scoring <- data.frame(AIC = max(-res.aic)/-res.aic, BIC = max(-res.bic)/-res.bic, MDL = max(-res.mdl)/-res.mdl, 1:4)

scoring.long <- melt(scoring, id.vars="X1.4")

ggplot(data = scoring.long, aes(x=X1.4, y=(value), group=variable, color=variable)) +
  geom_line() +
  geom_point() +
  ggtitle("Scoring in function of the number of children", subtitle = NULL) +
  xlab("# of parent per node") +
  ylab("% of max score") +
  scale_x_continuous(breaks=c(1,2,3,4,5,6,7))
```

# ASIA: HOW MANY PARENT ARE NEEDED?



# ASIA: CONSTRAINT-BASED LEARNING

```
##=====
## constraint-based algorithm
##=====

bn.gs <- gs(asia)
plot(bn.gs)

bn.iamb <- iamb(asia)
plot(bn.iamb)
```

