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Comparison of learning algorithms for Bayesian Networks models: a case study using the World Health Survey data

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BACKGROUND

- **AIM OF THE STUDY:** to compare algorithms for learning Bayesian Networks (BN) using the World Health Survey (WHS) data, a real dataset with numerous interdependences between variables.

WORLD HEALTH SURVEY (WHS):

- World Health Organization (WHO)
- 70 countries
- between 2002 and 2004
- cross-population comparable data on health, health-related outcomes and risk factors

WORLD HEALTH SURVEY

◦ **AIM OF THE WHS:** to provide valid, reliable and comparable information about the World population health status

SAMPLING DESIGN: probability sampling using multi-stage, stratified, random cluster samples

POPULATION STUDIED: persons aged 18 years and older who lived in households



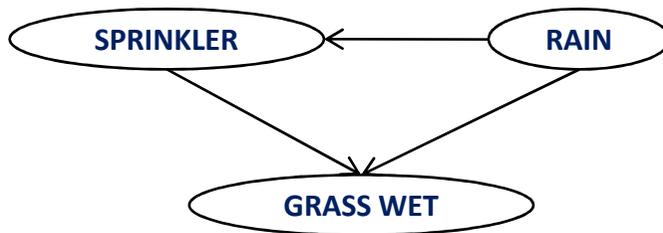
BAYESIAN NETWORK (1)

DEFINITION: BN is a Directed Acyclic Graph (DAG) whose nodes represent variables, and whose arcs describe the conditional in/dependencies between variables.

qualitative part
(BN structure)

quantitative part
(BN parameters)

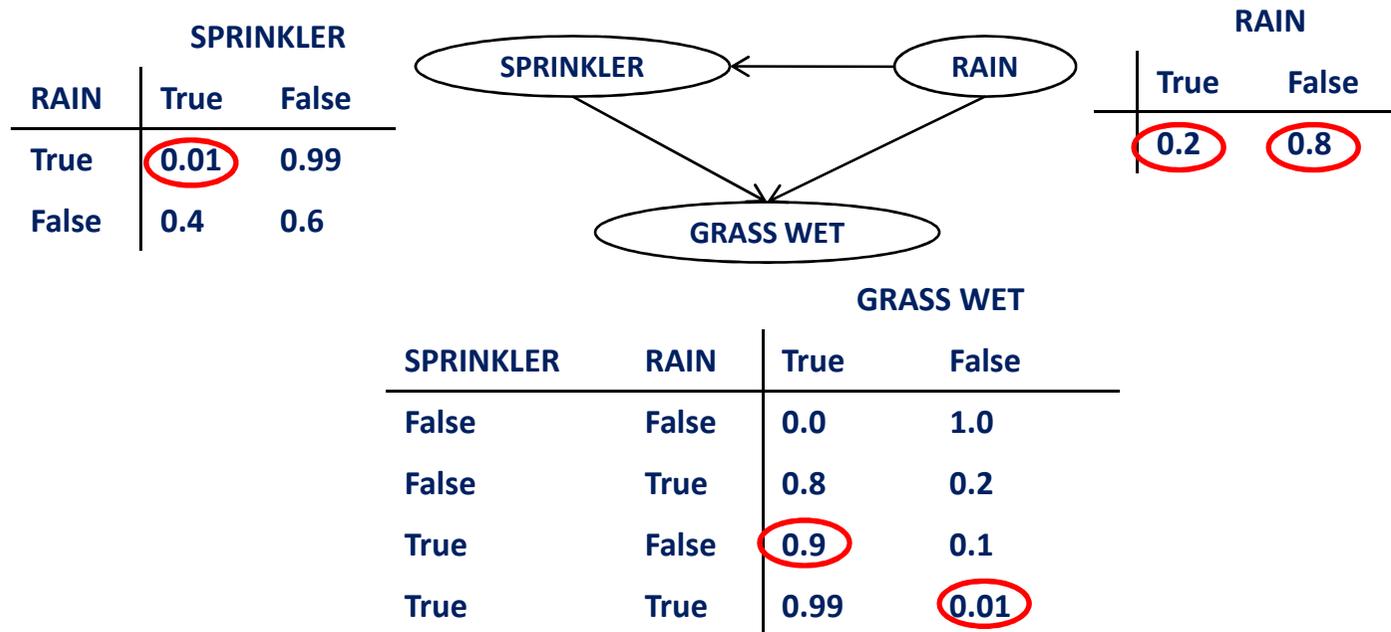
Example of the qualitative part



This example shows the structure of a BN. SPRINKLER is called a *child* of RAIN because the RAIN has a direct effect on the use of the SPRINKLER, and RAIN is called a *parent* of SPRINKLER.

BAYESIAN NETWORK (2)

Example of the quantitative part



This example shows the parameters of a BN.

The **joint probability function** of this BN is:

$$P(\text{RAIN}, \text{SPRINKLER}, \text{GRASS WET}) = P(\text{RAIN})P(\text{SPRINKLER} | \text{RAIN})P(\text{GRASS WET} | \text{SPRINKLER}, \text{RAIN})$$

BAYESIAN NETWORK (3)

BAYESIAN NETWORK is a graphical representation in the form of a DAG, G , for conditional in/dependencies and for compact specification of full joint distributions

G encodes the **Markov condition**: each node of the BN is probabilistically independent of its non-descendants, given its parents.

The **full joint distribution** is defined as the product of the local conditional distributions of each node of the network:

$$P(x_1, \dots, x_n) = \prod_{i=1}^n P(X_i \mid \text{Parents}(X_i))$$

BAYESIAN NETWORK APPLICATION (1)

Methods for construct a Bayesian network:

- (1) BN specified by an **expert**
- (2) BN learned from **data**

There are two primary approaches for learning a Bayesian Network from data:

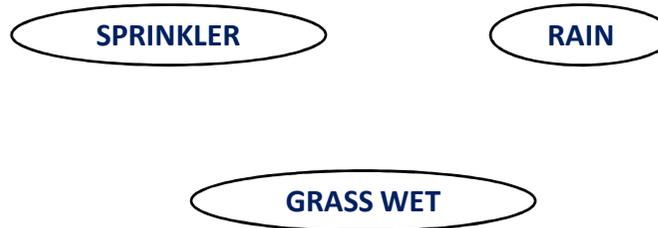
- (1) the constraint-based algorithms (**CI**), and
- (2) the search and score algorithms (**S&S**)

The **constraint-based** evaluates the presence or absence of an arc by testing conditional independencies.

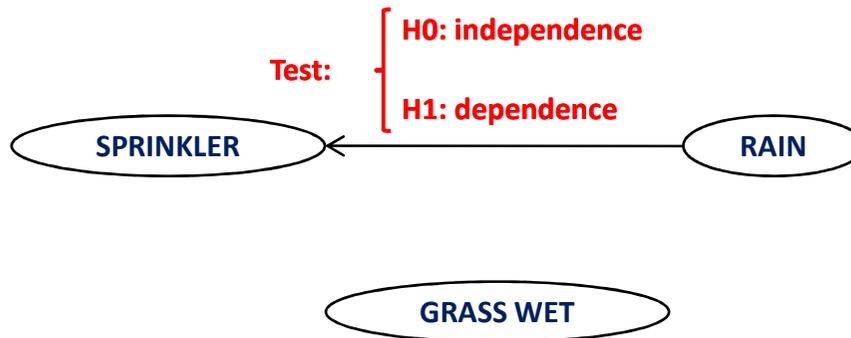
BAYESIAN NETWORK APPLICATION (1a)

Constraint-based method

1° step



2° step



BAYESIAN NETWORK APPLICATION (2)

- The **search and score** evaluates the goodness-of-fit of the network to the data maximizing a selected scoring function.

Typologies of **scoring functions** :

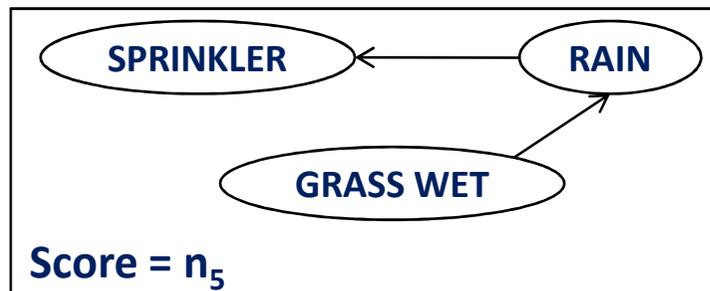
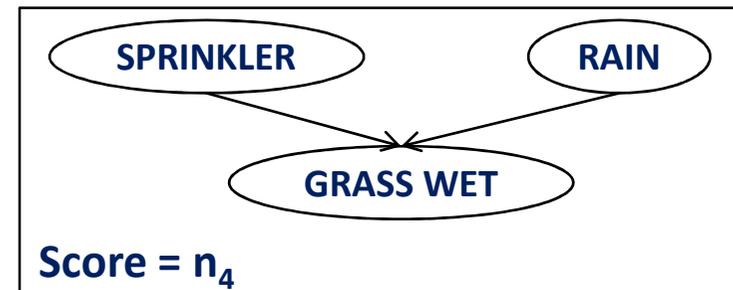
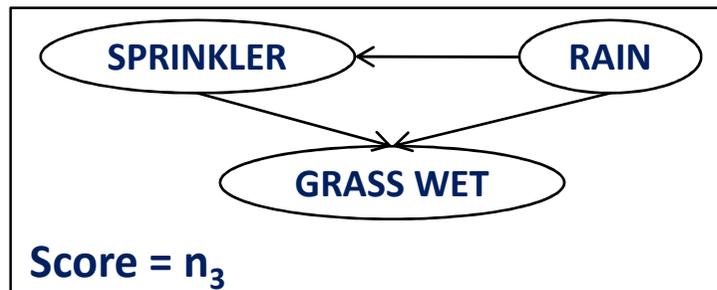
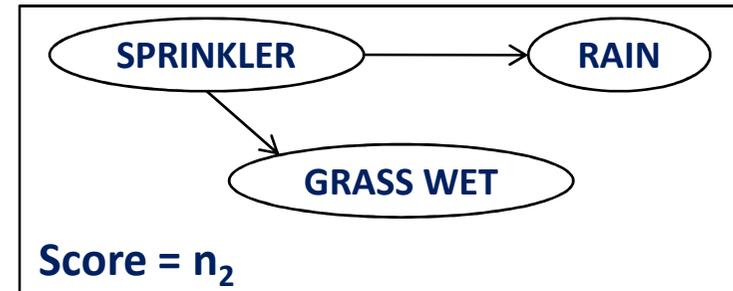
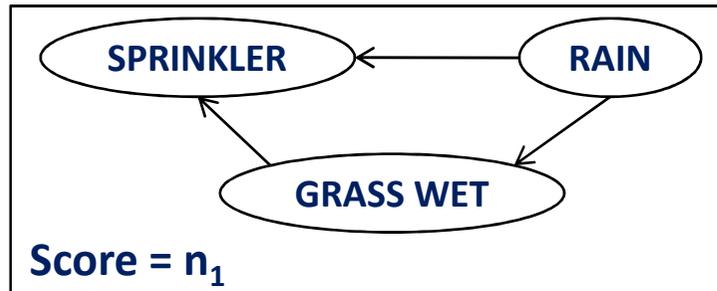
- (a) Bayesian (based on the Bayes theorem), and
- (b) Information Theory

The **Bayesian scoring functions** compute the posterior probability distribution conditioned to the data, starting from different prior probability distributions. The best network is the one that maximizes the posterior probability.

The **scoring functions based on Information Theory** select the network structure that best fits the data, penalized by the number of parameters of the network.

BAYESIAN NETWORK APPLICATION (2a)

Search and Score method



BAYESIAN NETWORK APPLICATION (3)

Constraint-based
approach



BNPC algorithm
(BNPC software)

Search & Score
approach



Tabu search algorithm
(Weka software)



Scoring function
BDeu
(Bayesian)



Scoring function
MDL
(Information Theory)



With uniform prior
distribution

CASE STUDY

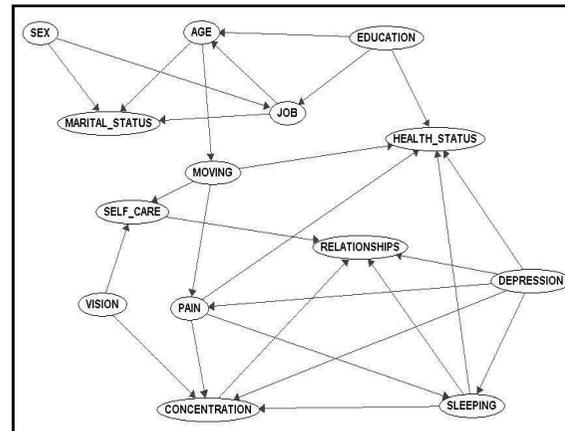
- The dataset used for the analysis contained **26,608** records from **22** countries.

14 categorical variables:

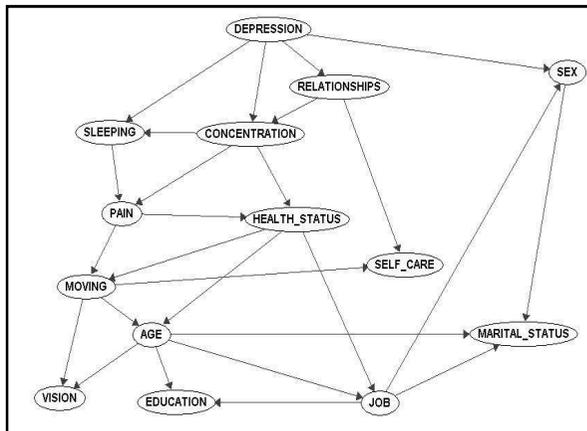
- **5** socio-demographic characteristics
(sex, age, marital status, education and employment)
- **1** self-reported overall health status question
- **difficulties in functioning in 8** health domains
(mobility, self-care, pain and discomfort, concentration, interpersonal relationships, vision, sleeping, and feeling sad or depressed)

RESULTS (1)

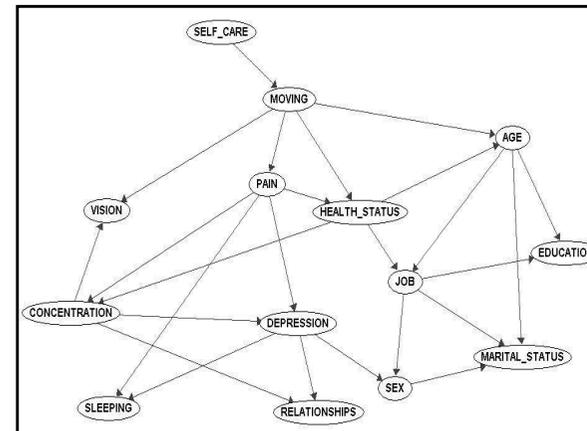
BNPC network (CI)



Tabu search network with scoring function BDeu (S&S)

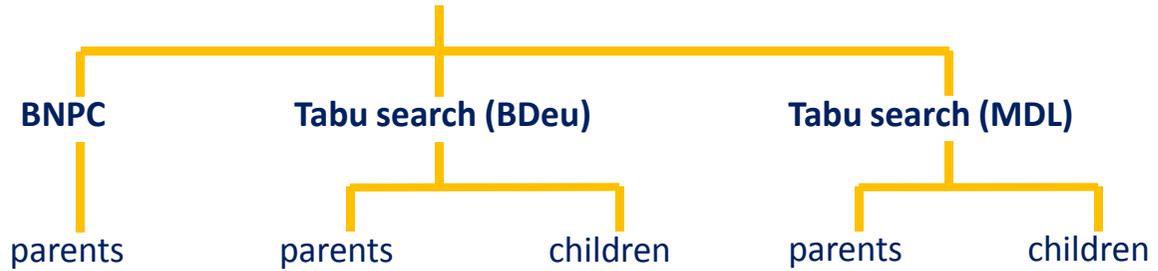


Tabu search network with scoring function MDL (S&S)



RESULTS (1a)

Health status



Socio-demographic domain

Psychological domain

Physical domain

Education

**Depression
Sleeping**

**Pain
Moving**

Concentration

Pain

**Age
Job**

Moving

**Age
Job**

**Pain
Moving**

Concentration

RESULTS (2)

° The principal aim of our study was to **compare** the different Bayesian networks, in terms of structure.

The comparison process is divided into two typologies:



The comparison of
the **structure** of
the network

The comparison of
the **predictive
accuracy** on the
target variable

RESULTS (3)

Comparison of the structure

	Tabu search (BDeu) vs Tabu search (MDL)	Tabu search (BDeu) vs BNPC	Tabu search (MDL) vs BNPC
Total number of arcs	26 vs 25	26 vs 27	25 vs 27
Coincident arcs*	15	8	11
Inverted arcs**	8	12	9
Added arcs***	2	7	7
Deleted arcs****	3	6	5

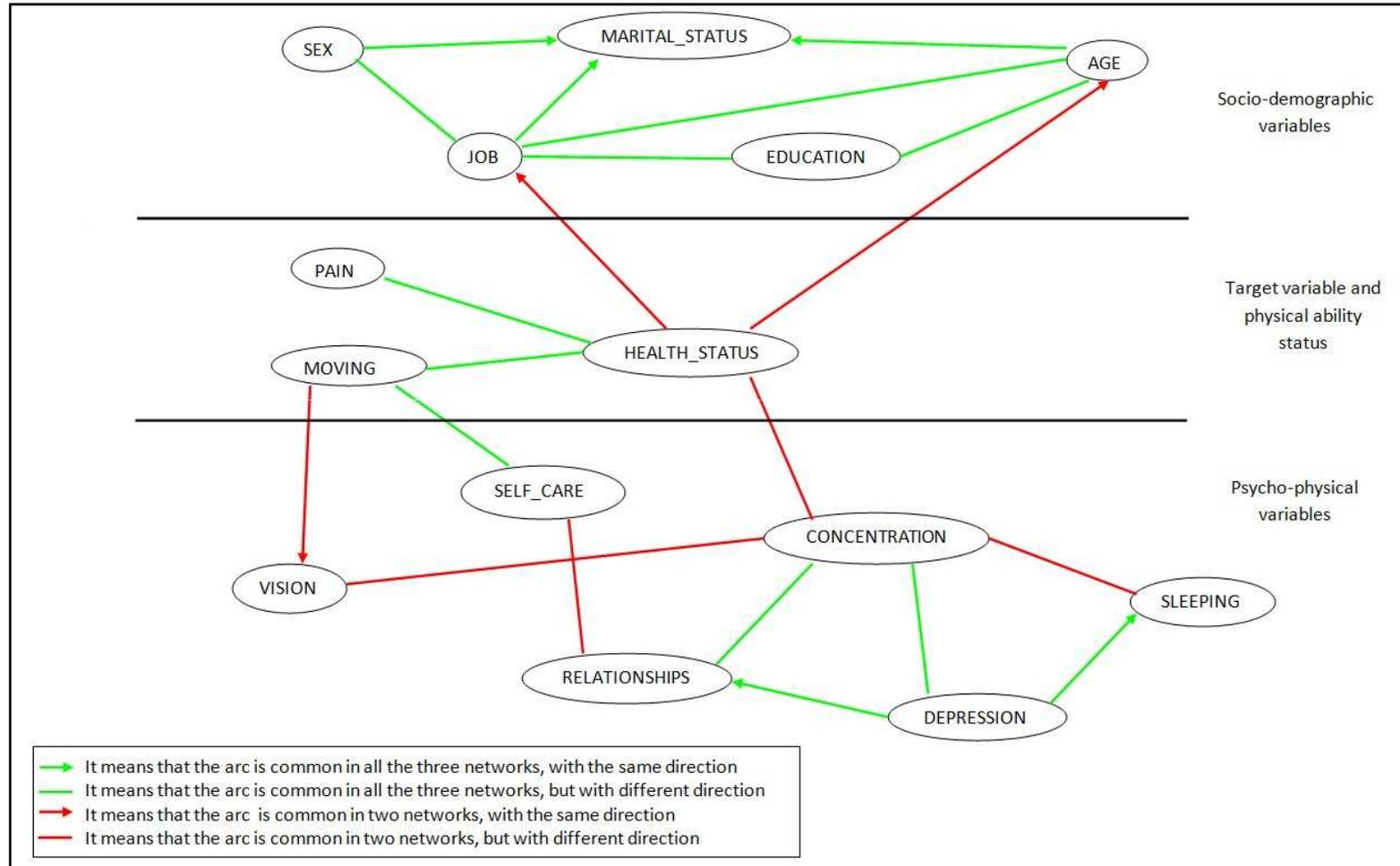
* An arc is coincident if it is present in both networks with the same direction

** An arc is inverted if it is present in both networks but with an inverted direction

*** An arc is added if it is not present in the first network but it is present in the second one

**** An arc is deleted if it is present in the first network but it is not present in the second one

RESULTS (3a)



RESULTS (4)

Comparison of the predictive accuracy

Each BN was used to predict the most probable value of the health status variable and the comparison of the predicted with observed values produced the **percentage of classification success**

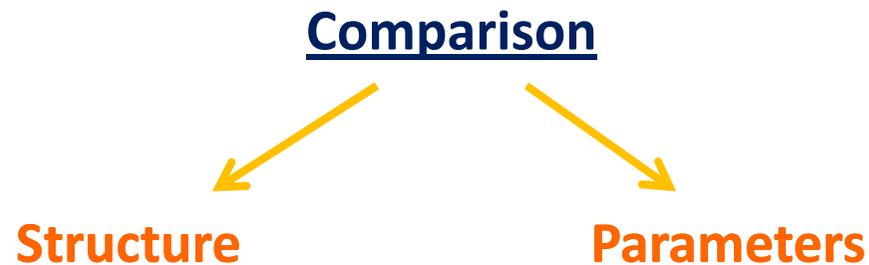
	Tabu search (BDeu)	Tabu search (MDL)	BNPC*
Percentage of classification success (SD)	51.77% (0.51)	51.41% (0.52)	49.72% (na**))

* The BNPC software does not support the calculation of the Standard Deviation

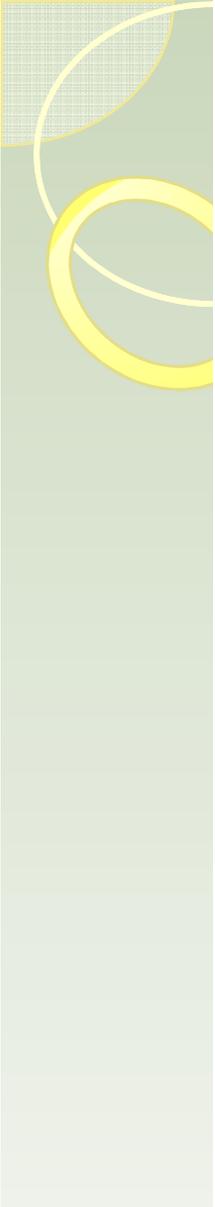
** Percentage of classification success was performed with HUGIN software

CONCLUSIONS

- We have compared two different typologies of algorithms, which are based on different assumptions



- **Strength:** high dimension WHS dataset
- **Limitation:** different characteristics of the software used



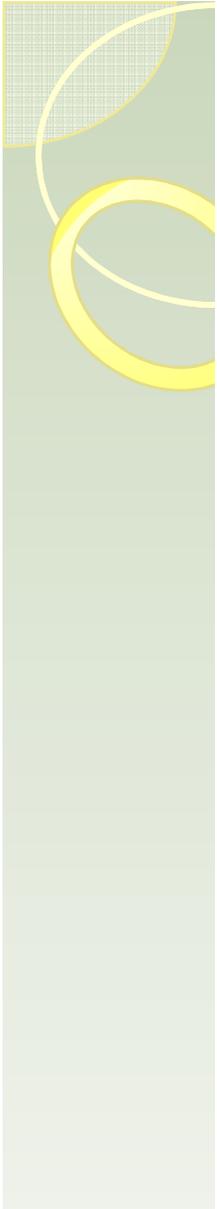
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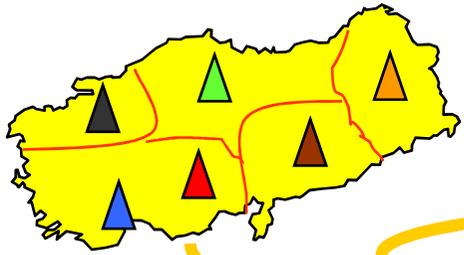
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Thank you very much for your attention!







**Provinces
Strata**



100 Counties - PSUs



Enumeration Areas - SSUs



50 Households - TSUs



Respondents

Bayesian scoring functions (based on Bayes theorem)

$$P(G|D) \sim P(G) P(D|G)$$

↓ ↓ ↓

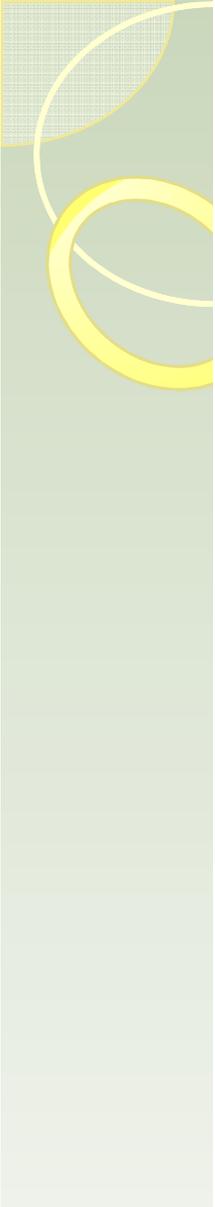
posterior prior likelihood

Scoring functions based on Information theory

$$MDL(G|D) = LL_D(G) - \frac{1}{2} C(G) \log(N)$$

↓ ↓ ↓

score log-likelihood network complexity



K-fold cross validation

The overall dataset is randomly partitioned into 10 subsets of approximately equal size. Of the 10 subsamples, a single subsample is retained as the validation data for testing the model (the test set), and the remaining nine subsamples are used as training data (the training set). This process is then repeated 10 times.

Simple cross validation

The overall dataset is split into two subsets: one training set and one test set.