

**gRaphical models:
a softwaRe peRspective**

Steffen L. Lauritzen, Aalborg University

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Why this lecture?

- Graphical models have been around for about 25 years
- Software is the most important vehicle for dissemination of statistical ideas into practice
- Graphical models have shown some potential
- Software for graphical models exists as several independent stand-alone packages
- Time has come to attempt integration into general, flexible, and extendable software, such as R; hence **gR**.

Overview

- Brief historical sketch
- Basics of graphical models
- Present software situation
- Key dilemmas and challenges
- Basic needs and easy wins
- Demanding abstractions
- Where to go from here

Undirected graphical models

- Traces back to statistical physics (Gibbs 1902)
- Models for spatial interaction (Besag 1974)
- Interpreting hierarchical log-linear models by conditional independence, using analogy to Markov random fields (Darroch, Lauritzen and Speed 1980); **CoCo** (Badsberg 1991)
- Extending hierarchical log-linear models to include continuous variables (Lauritzen and Wermuth 1989); **MIM** (Edwards 1990).

Directed graphical models

- Traces back to path analysis (Wright 1921)
- Generalizing (block) recursive linear systems to discrete case (Wermuth and Lauritzen 1983, Wermuth and Lauritzen 1990); **DIGRAM** (Kreiner 1989).
- Bayesian (causal) networks (Pearl 1986, Lauritzen and Spiegelhalter 1988); **HUGIN** (Andersen, Olesen, Jensen and Jensen 1989), **TETRAD** (Spirtes, Glymour and Scheines 1993). **DEAL** (Bøttcher and Dethlefsen, DSC 2003).
- Bayesian graphical modelling; **BUGS** (Gilks, Thomas and Spiegelhalter 1994).

Undirected graphical models

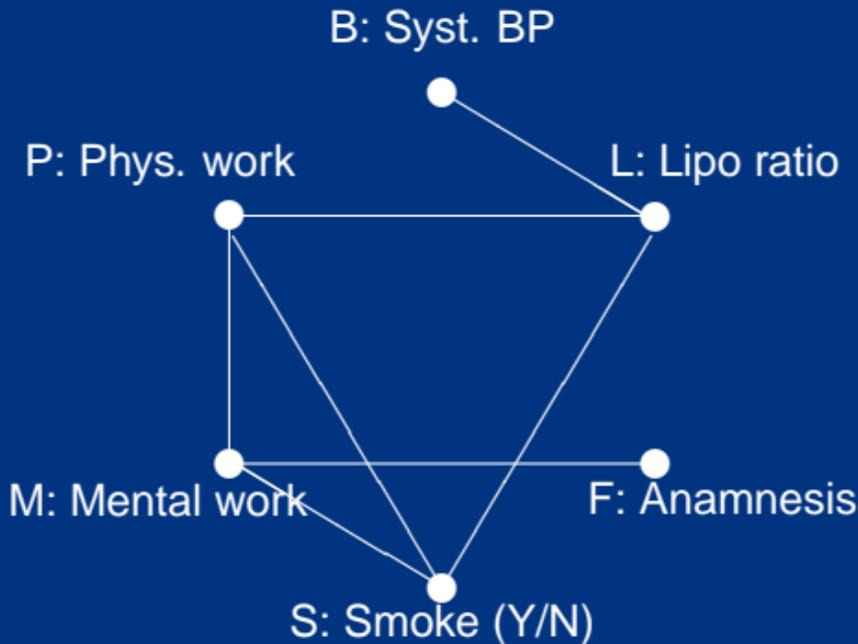
- Nodes V represent set of variables $X_v, v \in V$
- Undirected graph $\mathcal{G} = (V, E)$ with (maximal) cliques \mathcal{C} .
- Joint density factorizes as

$$f(x) = \prod_{c \in \mathcal{C}} \phi_c(x),$$

where ϕ_c depends on x through $x_c = (x_v)_{v \in c}$ only.

- Conditional independence: $A \perp\!\!\!\perp B \mid S$ if set of variables S separates A from B in \mathcal{G} .

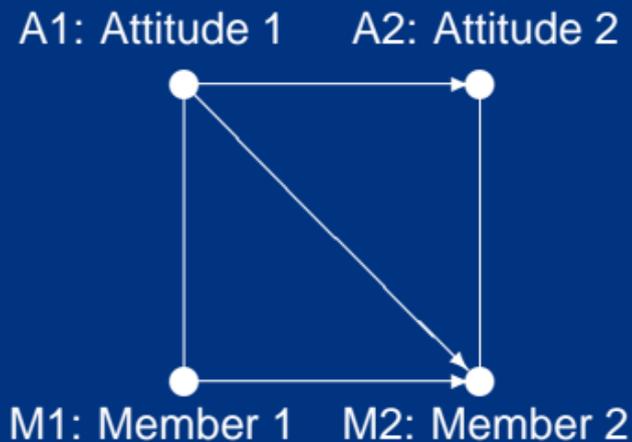
Undirected graphical models



$F \perp\!\!\!\perp S, B, P, L \mid M$ and $F, M \perp\!\!\!\perp L, B \mid S, P$ and
 $B \perp\!\!\!\perp S, P, M, F \mid L$.

show MIM

Directed (chain) graphical models



$$A2 \perp\!\!\!\perp M1 \mid A1, M2$$

show DIGRAM

Models for data

- Models shown so far are **models for data**.
- Graph represents **model formula**
- Parameters are **implicit** and not part of graph
- Data themselves are separate and **not** represented in graph
- Models are in essence **multivariate**
- Integration into R is 'straightforward' and fits with usual paradigm

Some simple advances

- Make interfaces between e.g. MIM and R (Højsgaard, DSC 2003)
- Make e.g. CoCo available within R (Badsberg, DSC 2003)
- Make R packages for special purposes e.g. DEAL (Bøttcher and Dethlefsen, DSC 2003)
- Make facilities directly within R so graphical modelling becomes easily extendable.

gRaphical models for data

Facilities to be desired within R to perform relevant analyses:

- Facilities for manipulating, representing, interacting with, displaying and printing graphs
- facilities and algorithms for specifying fitting, testing, selecting graphical models as in stand-alone programs
- Facilities for exporting and importing models and analyses to other programs

Bayesian networks

- Directed Acyclic Graph \mathcal{D} (DAG)
- Nodes V represent variables $X_v, v \in V$
- Specify conditional distributions of children given parents: $p(x_v | x_{\text{pa}(v)})$
- Joint distribution is then $p(x) = \prod_{v \in V} p(x_v | x_{\text{pa}(v)})$
- Algorithm transforms network into *junction tree*
- **Inference** is performed by computing $p(x_v | x_A)$ which can be efficiently computed for all $A \subseteq V$.

show HUGIN

Data for models

BUGS extends idea of Bayesian networks for statistical purposes (complex Bayesian modelling)

- Parameters **explicit**, represented as nodes in graph
- Data explicitly represented in graph by **observational nodes**
- Special symbolism for **repeated structures** (plates)
- Inference by updating from prior to posterior, in effect **calculating likelihoods** by MCMC.

show BUGS

Compare simple linear regression

- Classical graphical model



- Standard model formula in R:

```
lm(y~x, data)
```

- BUGS model specification:

```
model{for(i in 1:N)
  {Y[i]~ dnorm(mu[i], tau)
  mu[i] <- a + b * (x[i] - mean(x[]))}
}
```

Contrasting paradigms

Bayesian vs. classical **inference** are not very different:

‘Classical’ graphical models can be treated with Bayesian methods

- Specify prior distributions
- Integrate likelihood functions rather than maximize
- Bayes factors instead of significance tests
- Posterior model probabilities for model search rather than comparison with significance tests, etc.

Asymptotically indistinguishable

Contrasting paradigms

Differences more radical concerning **attitude towards modelling**

- **Data for models** rather than models for data
- parameters, data, latent variables, covariates are **fundamentally the same**, i.e. random variables
- Status of variables changes **dynamically** through observation: X becomes x when observing $X = x$.
- Builds complex models using **modularity** and **local modelling**

Towards computing with models

- Local modelling paradigm is **not** Bayesian, but more fundamental
- Fits with **modularity** of graphical models
- In principle one could make LUGS, a likelihood version of BUGS. Just use prior distributions as 'convenient computational device'.
- More **difficult to integrate** local modelling paradigm into R. R has its origin within 'data analysis'; 'models' a later 'add on'.
- Needs **further abstraction** of data structures and object orientation.

Heads and Tails

Need **local model object** (LMO) $cd(H | T)$ with **head** H and **tail** T .

Heads correspond to random variables, tails to parameters, in traditional thinking.

Variables in heads and tails can be **instantiated** by data. When T is empty or fully instantiated, $cd(H | T)$ is a **distribution**, possibly represented as a program which simulates from it or otherwise can integrate.

When H is fully instantiated, $cd(H | T)$ is a **likelihood** and is represented as a function which can be evaluated.

Operations with Models

With heads partially instantiated, LMOs represent **conditional distribution** of uninstantiated part of head given tail and instantiated part of head.

LMOs can be **marginalised** over uninstantiated parts of head. This reduces their head.

LMOs $cd(H_1 | T_1)$ and $cd(H_2 | T_2)$ can be **combined** when either $H_1 \cap (H_2 \cap T_2) = \emptyset$ or $H_2 \cap (H_1 \cap T_1) = \emptyset$ (or they satisfy specific consistency condition).

Graphs are formal objects which keep track of how LMOs are combined.

Where to go from here

- Continue first integration of stand-alones into R
- Identify specific elements of stand-alones which could usefully become part of gRaphical model library
- Identify new useful features which could usefully be added to such a graphical model library
- make repository of data for graphical models
- Do it!
- Continue process of abstraction to realise 'computation with models'

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