

# Abstracting Complex Systems using Mixed Graphical Models

Jonas Haslbeck

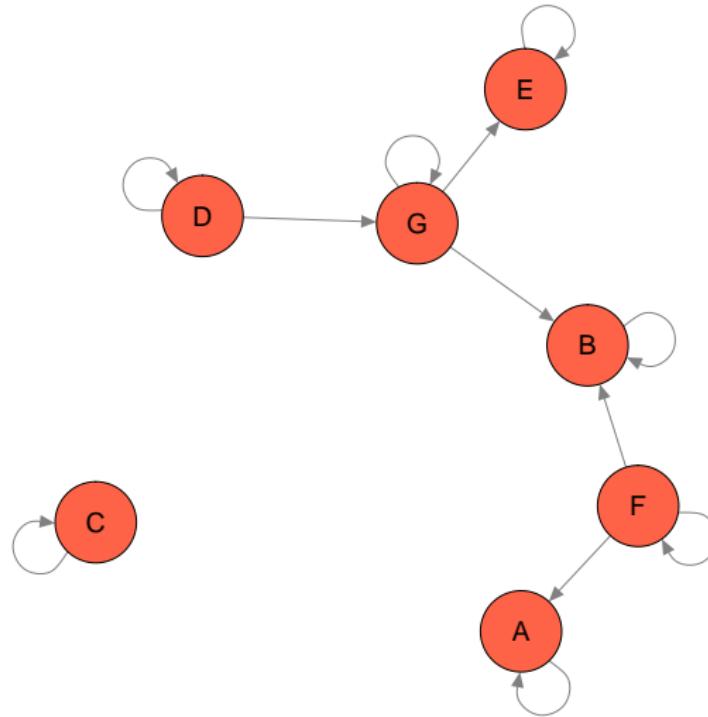
*Psychosystems lab  
University of Amsterdam, the Netherlands*

[psychosystems.org](http://psychosystems.org)  
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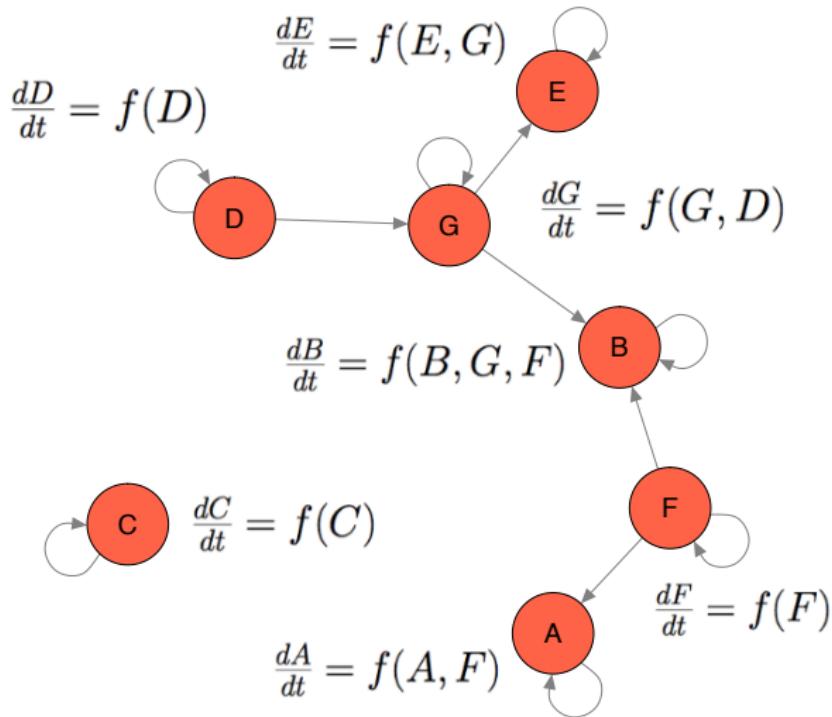
Complexity Laboratory Utrecht (CLUe) Lunch Meeting

Utrecht, October 20th

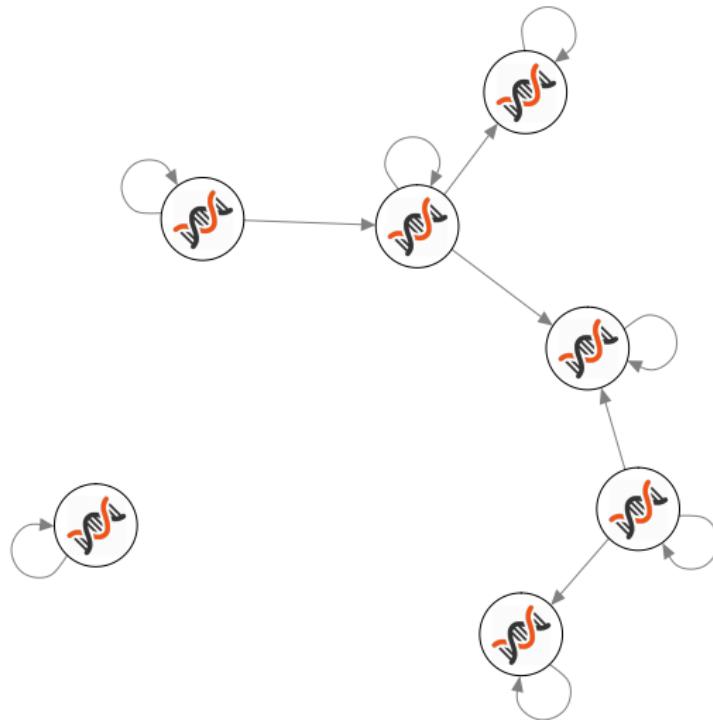
# Multivariate System



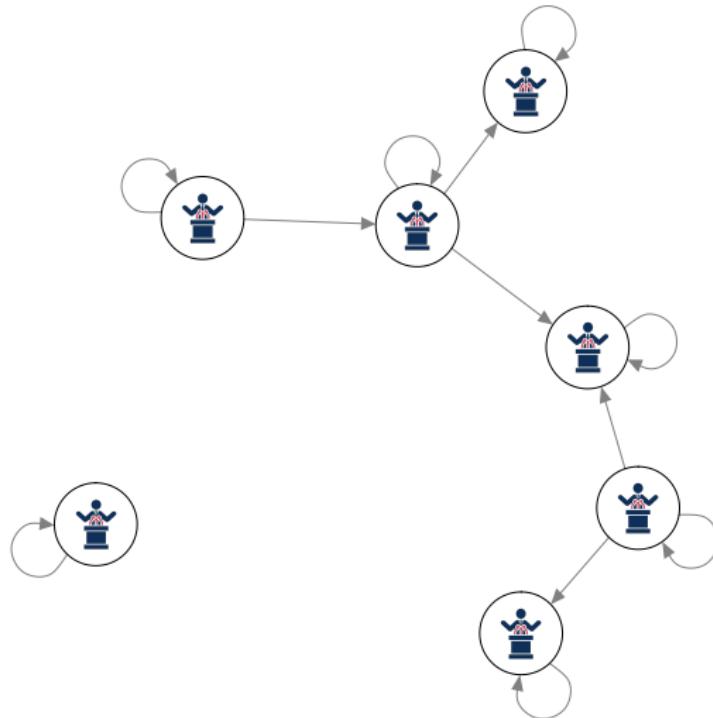
# Multivariate System



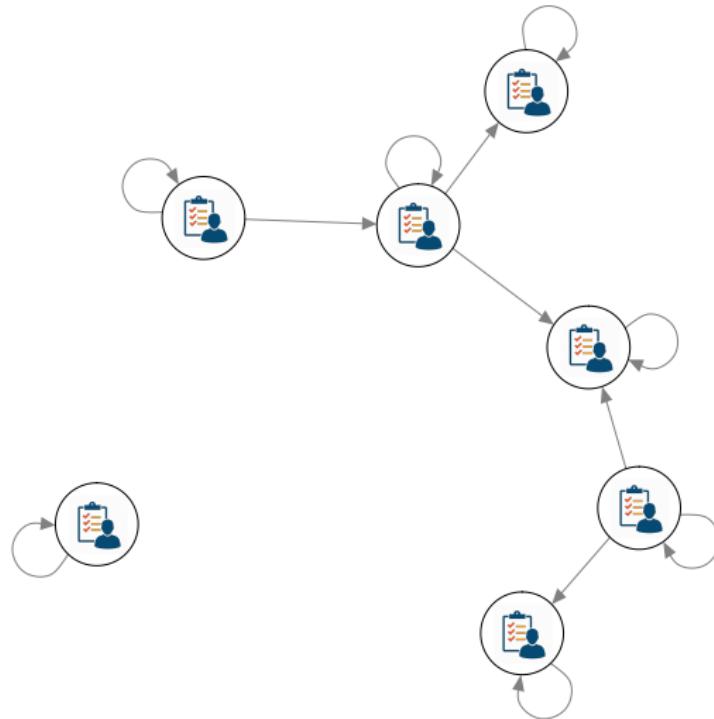
# Gene Expressions



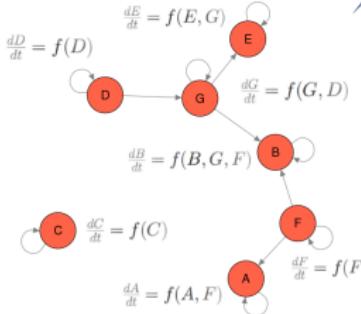
# Voting Behavior of Members of Parliament



## Symptoms of Mental Disorders



## Sample observations

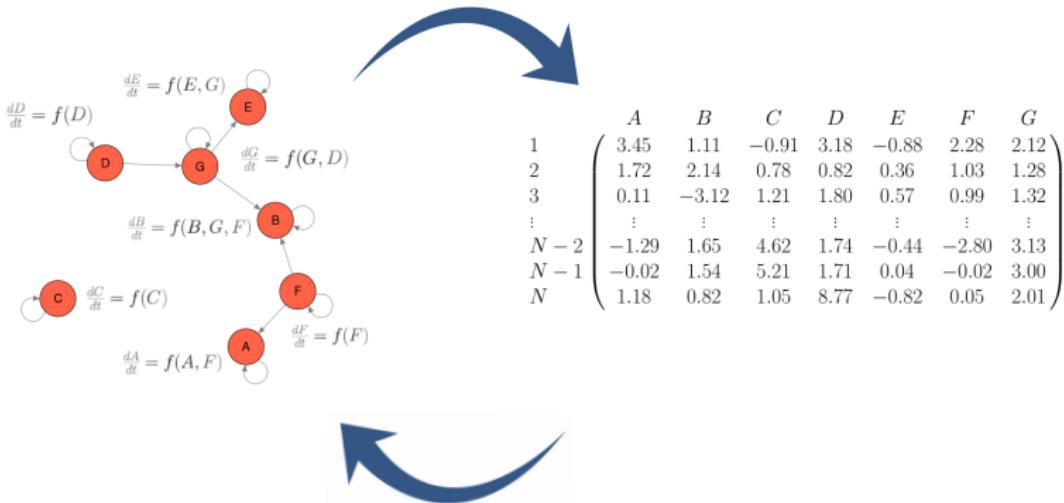


	$A$	$B$	$C$	$D$	$E$	$F$	$G$
1	3.45	1.11	-0.91	3.18	-0.88	2.28	2.12
2	1.72	2.14	0.78	0.82	0.36	1.03	1.28
3	0.11	-3.12	1.21	1.80	0.57	0.99	1.32
:	:	:	:	:	:	:	:
$N - 2$	-1.29	1.65	4.62	1.74	-0.44	-2.80	3.13
$N - 1$	-0.02	1.54	5.21	1.71	0.04	-0.02	3.00
$N$	1.18	0.82	1.05	8.77	-0.82	0.05	2.01



Recover the system

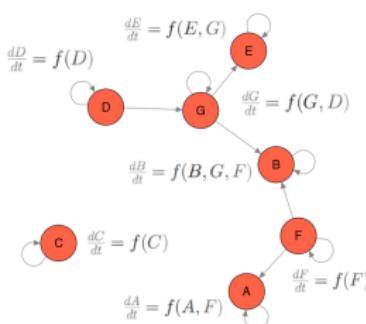
## Sample observations



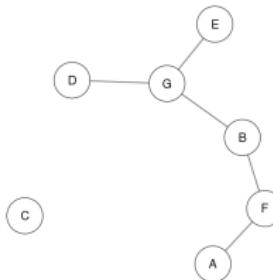
Approximate the system

# True Model, Probability Distribution, Network Model

True Model



Conditional Independence Network



Approximate

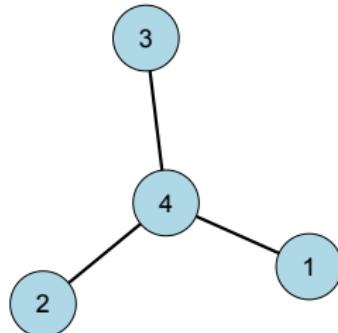
Summarize

$$P(X_1, \dots, X_p, \theta)$$

Multivariate Probability Distribution

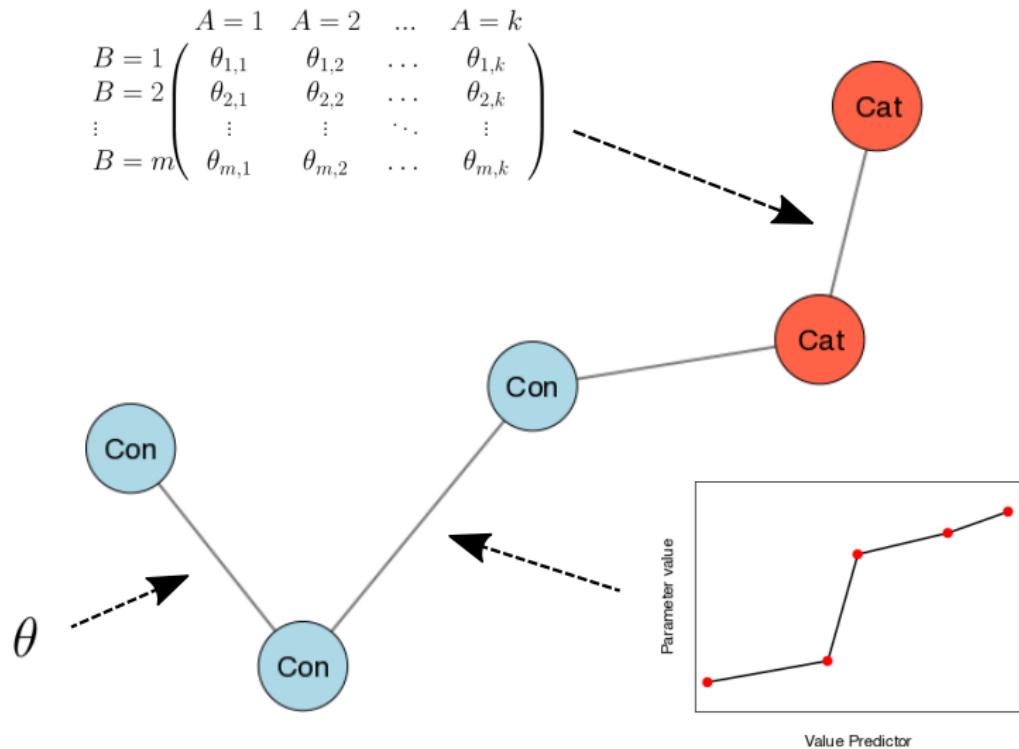
## Simple Example: Gaussian Graphical Model

$$\Sigma^{-1} = \begin{matrix} & \begin{matrix} X_1 & X_2 & X_3 & X_4 \end{matrix} \\ \begin{matrix} X_1 \\ X_2 \\ X_3 \\ X_4 \end{matrix} & \begin{pmatrix} 3.45 & 0 & 0 & 3.18 \\ 0 & 2.14 & 0 & 0.82 \\ 0 & 0 & 3.21 & 1.05 \\ 3.18 & 0.82 & 1.05 & 8.77 \end{pmatrix} \end{matrix} \iff$$



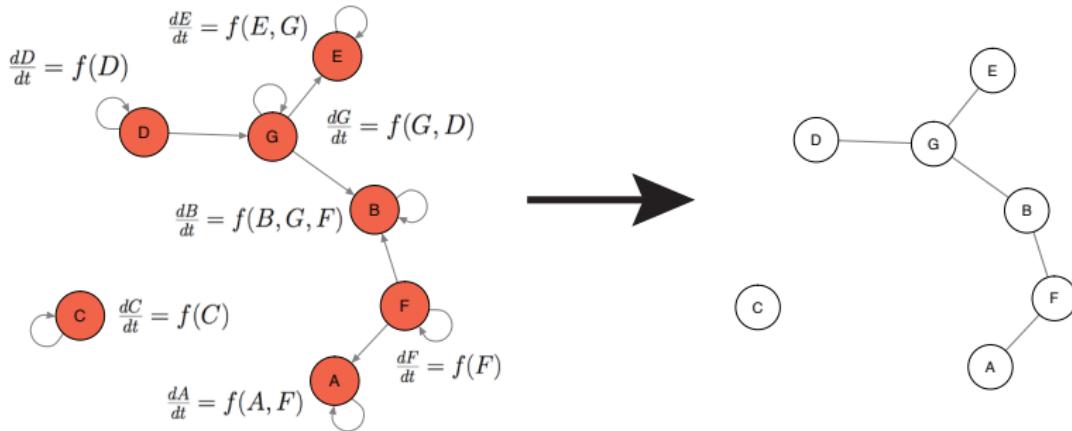
$$P(X_1, \dots, X_p) = \frac{1}{\sqrt{(2\pi)^p |\Sigma|}} \exp \left\{ -\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu})^\top \Sigma^{-1} (\mathbf{x} - \boldsymbol{\mu}) \right\}$$

# General Graphical Models



## Goal:

Abstract structure of true system in simpler MGM model class



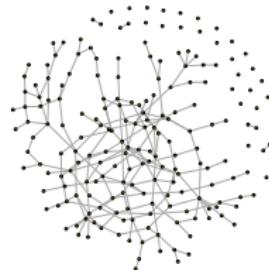
True System

Abstraction in simpler  
MGM model class

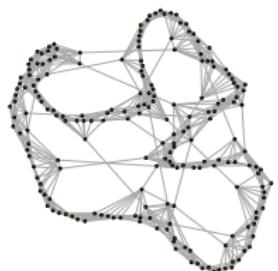
# Study multivariate distribution as network



Ring Network



Random Network

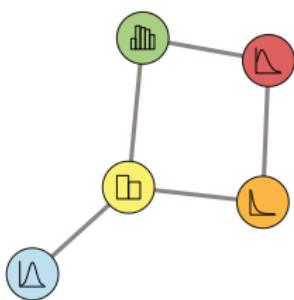


Small World Network

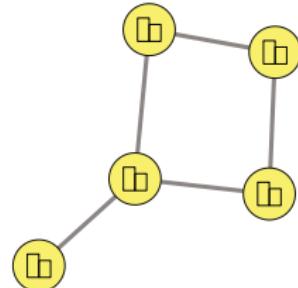


Scale-free Network

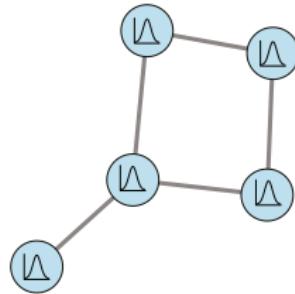
# Mixed Graphical Models



Mixed Graphical Model



Ising Model



Gaussian Graphical Model

# Constructing MGMs

Each node/variable is a univariate exponential family distribution conditional on all other variables

$$P(X, \beta) = \exp \{ \eta(\theta)B(X) + C(X) - A(\theta) \}$$

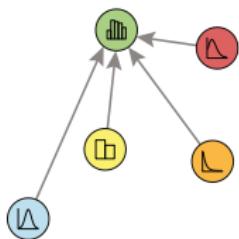
and the natural parameter  $\theta$  is a linear combination of all other variables:

$$\theta_{t,i} = \beta_{0,i} + [\beta_{i,1} \ \dots \ \beta_{i,p}] \begin{bmatrix} X_2 \\ \vdots \\ X_p \end{bmatrix}$$

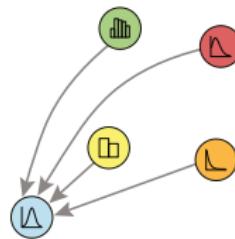
(Yang et al., 2014; Chen, Witten & Shojaie, 2015; Haslbeck & Waldorp, 2017)

# Estimating Mixed Graphical Models

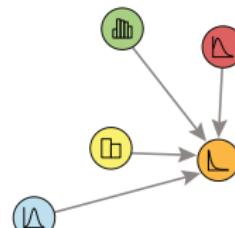
Step 1



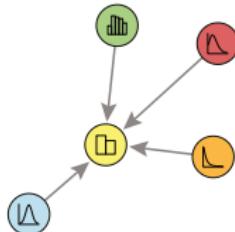
Step 2



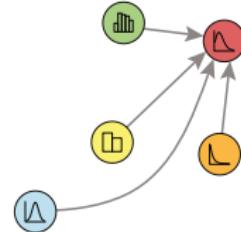
Step 4



Step 3



Step 5



(Meinshausen & Bühlmann, 2006)

## $\ell_1$ -regularized Estimation

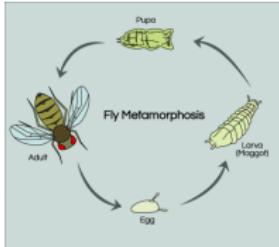
We minimize the negative log-likelihood  $F(\mathbf{X}_j, \boldsymbol{\theta}_t)$  together with the  $\ell_1$ -norm over all parameters:

$$\arg_{\boldsymbol{\theta}_t} \min \left\{ \frac{1}{n} \sum_{j=1}^n w_{j,t_e} F(\mathbf{X}_j, \boldsymbol{\theta}_t) + \lambda_i \|\boldsymbol{\theta}_t\|_1 \right\}$$

This has two consequences:

1. We can control the bias (model too simple) vs. variance (model too complicated) trade-off with tuning parameter  $\lambda$ ;
2. Small parameters are set to exactly zero

# Back to Applications



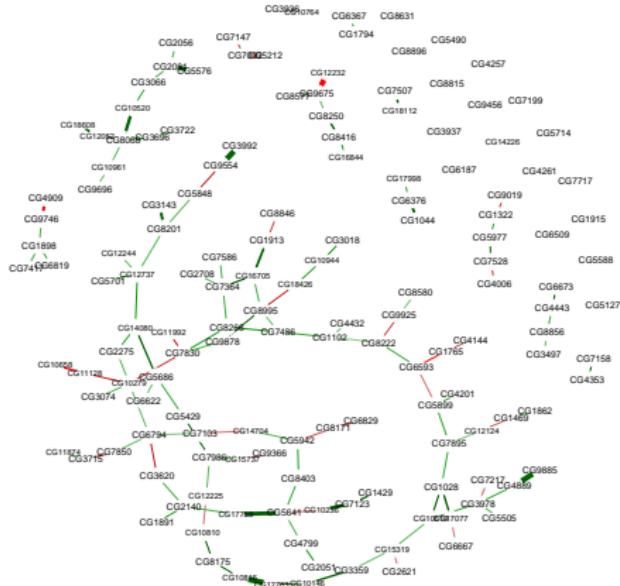
67 measurements of 150 gene expressions related to the immune system of *Drosophila melanogaster* (fruit fly) over its full life cycle

Votes of 623 members of the German parliament on 136 bills from Nov 2013 - April 2015



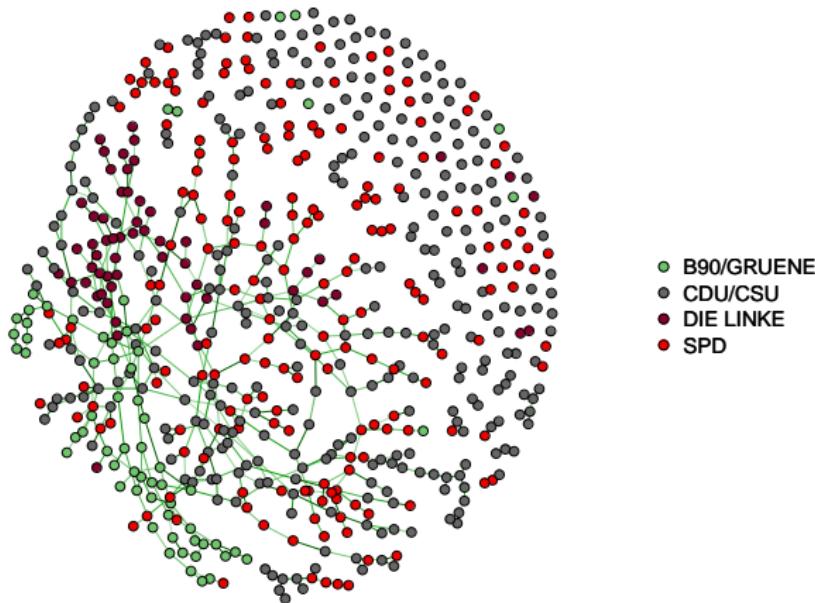
1476 measurements of 16 mood related variables of one individual over 238 consecutive days

# Gene Expressions of Fruit Fly



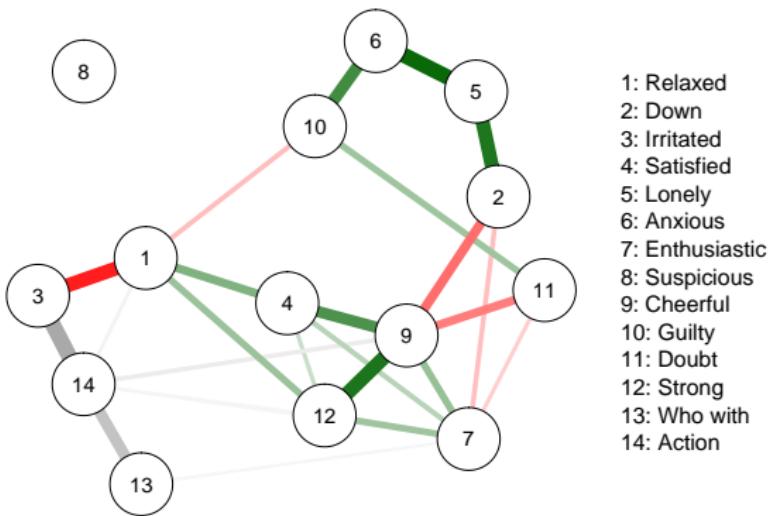
67 Measurements of 150 genes expressions related to immune system of the fruit fly (Lebre et al., 2010)

# Voting Behavior of Members of German Parliament



136 public votes, 623 members of parliament of 4 parties

# Symptoms of Mental Disorder



1476 measurements of 14 variables related to mood, activity and social context of one individual over 238 consecutive days  
(Kossakowski et al., 2017)

Practical:

Estimate MGM on Symptom Data

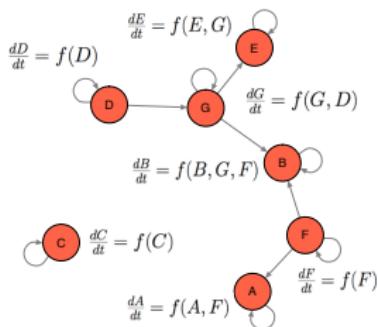
**RStudio Server:**

<http://clue.science.uu.nl:8787>

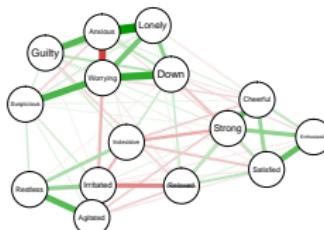
Login: Your UU Solis-ID & password

# Direction of Influence & Interactions as function of time

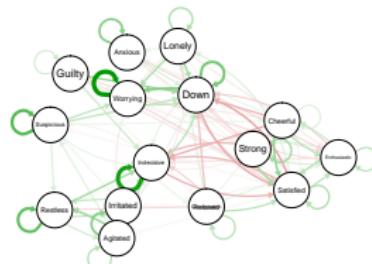
True Structure:



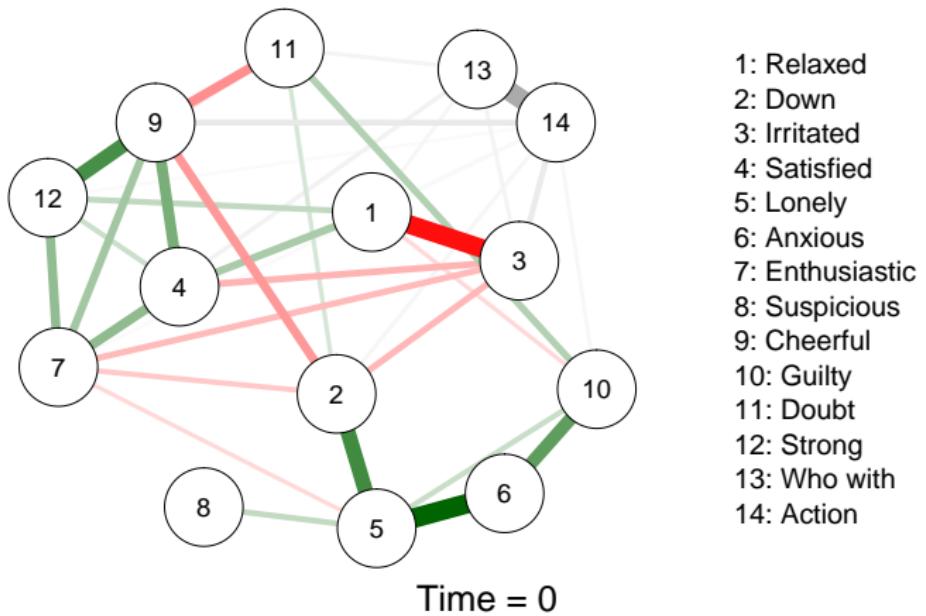
Instantaneous Influence



Influence over time (1h)



Does the system under investigation change over time?

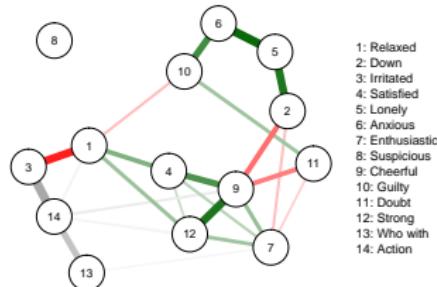


Does the system under investigation change over time?

# *mgm*: Summary

*mgm* package implements:

- ▶ Mixed Graphical Models (MGMs)
- ▶ Time-varying MGMs
- ▶ mixed Vector Autoregressive (mVAR) models
- ▶ Time-varying mVARs



**Website:** [jmbh.github.io](https://jmbh.github.io)

**Email:** [jonashaslbeck@gmail.com](mailto:jonashaslbeck@gmail.com)

## References

- ▶ Haslbeck, J., & Waldorp, L. J. (2017). Estimating mixed graphical models in high-dimensional data. arXiv preprint arXiv:1510.05677.
- ▶ Yang, E., Baker, Y., Ravikumar, P., Allen, G., & Liu, Z. (2014, April). Mixed graphical models via exponential families. In Artificial Intelligence and Statistics (pp. 1042-1050).
- ▶ Chen, S., Witten, D. M., & Shojaie, A. (2014). Selection and estimation for mixed graphical models. *Biometrika*, 102(1), 47-64.
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- ▶ Lebre, S., Becq, J., Devaux, F., Stumpf, M. P., & Lelandais, G. (2010). Statistical inference of the time-varying structure of gene-regulation networks. *BMC systems biology*, 4(1), 130.
- ▶ Meinshausen, N., & Bhlmann, P. (2006). High-dimensional graphs and variable selection with the lasso. *The Annals of Statistics*, 1436-1462.