

Modeling Intensive Longitudinal Data in Discrete and Continuous Time

R group University of Zurich
Day 1

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Postdoc at the Department of Methodology & Statistics,
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Psychology (Bachelors) and Methods & Stats (Masters,
PhD)

Originally from Ireland (hence, “Oisín” → [Uh-sheen])

Research focuses on Dynamical Systems and Causal
Modeling in (Clinical) Psychology

PhD supervisor: Ellen Hamaker



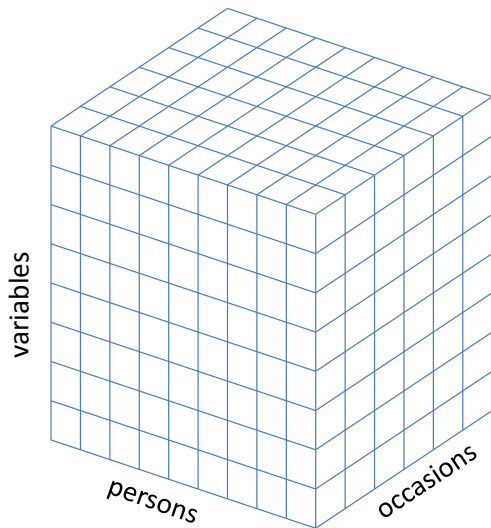
Universiteit Utrecht

Aim: Introduce you to the basic principles in modeling *intensive longitudinal data*

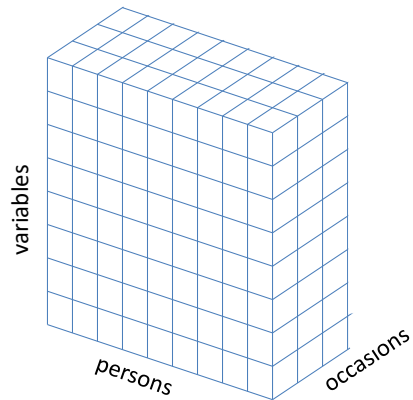
Broadly:

- Two related frameworks: Discrete-Time and Continuous-Time modeling
- Both model change over time using longitudinal data / repeated measures of the same variables
- Models built on *lagged regression*: X now predicts X later
- Can be viewed through the lens of SEM
- Primarily *single subject* approaches which can be generalized to a *multi-level* setting
- Focus on conceptual understanding with practical parts in *R*

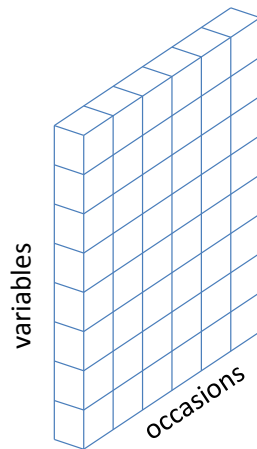
Cattell's data box



Panel Data: N = large and T is small (e.g. 2 -5 measurements)



Intensive Longitudinal Data: $N=1$ or many, T is large



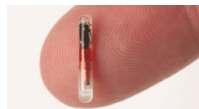
- Single ($N=1$) or multiple subjects ($N = \text{many}$). Many repeated measurements
- Time-points spaced closely together
- Attempts to directly capture short time-scale processes
- Focus primarily on individual within-person (idiographic) structure
- Sometimes called “time series” data
- “Long” format data

New technology

Smart phones

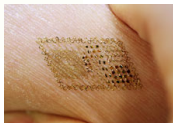


Smart glasses



Implants

Smart tattoo



Secure
continuous
remote alcohol
monitor
(SCRAM)



Smart watches



Activity trackers



Different forms of intensive longitudinal data:

- daily diary (DD); self-report end-of-day
- experience sampling method (ESM); self-report of subjective experience
- ecological momentary assessment (EMA); healthcare related self-report
- ambulatory assessment (AA); physiological measurements
- event-based measurements; self-report after a particular event
- observational measurements; expert rater

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For more info on **methodology**, check out:

- Seminar of Tamlin Conner and Joshua Smyth on YouTube (<https://www.youtube.com/watch?v=nQBBVp9vBIQ>)
- Society for Ambulatory Assessment (<http://www.saa2009.org/>)
- Life Data (<https://www.lifedatacorp.com/>)
- Quantified Self (<http://quantifiedself.com/>)

Characteristics of these kind of data

Data structure:

- one or more measurements per day
- typically for multiple days
- sometimes multiple waves (i.e., Nesselroade's measurement-burst design)

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Proposed Advantages of ESM, EMA and AA

- no recall bias (in comparison to retrospective questionnaires)
- high ecological validity (in comparison to lab experiments)
- physiological measures over a large time span
- monitoring of symptoms and behavior, with new possibilities for feedback and intervention (e-Health and m-Health)
- window into the dynamics of processes

Discrete and Continuous-Time Models

Discrete-Time (DT) models

- Current observation regressed on previous observation(s)
- *Time series analysis* tradition of economics

Continuous-Time models

- Based on *differential equations*: instantaneous change as a function of current value
- *Dynamical Systems Theory* - Physics, Ecology, Biology

Different perspectives/emphases, but many shared concepts and connections

- Qualitative behaviour of the systems being modeled!

- **Time series analysis: Univariate**
- Time series analysis: Multivariate
 - Practical: Exercises 1, 2 and 3
- Multilevel time series analysis
 - Practical: Exercise 4 & 5
- Extensions and Advanced Issues
- Discussion

What is time series analysis?

Time series analysis is a class of techniques that is used in econometrics, seismology, meteorology, control engineering, and signal processing.

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Main characteristics:

- $N=1$ technique
- T is large (say >50)
- concerned with *trends*, *cycles* and *autocorrelation structure* (i.e., **lagged** relationships)
- traditional emphasis on models for *forecasting*

TSA in the social and medical sciences

In **sociology**:

- quarterly unemployment numbers
- effect of alcohol consumption per capita on criminal violence rates
- effect of suicide news on suicide rates

In **medical research**:

- effect of safety warnings on antidepressants use
- effects of pain control strategies
- effect of 9/11 attacks on weekly psychiatric patient admissions

In **psychology**:

- network of symptoms in depressive patient
- effect of feedback on academic performance
- effect of an intervention on the relationship between stress and affect

Time Series Analysis

Time-Series Models typically take the form

$$Y_t = f_1(T) + f_2(S) + f_3(Y_{t-1}, \dots, Y_{t-s}) + \varepsilon_t$$

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- A random / noise / residual / innovation term ε_t

Time Series Analysis

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Focus: picking an appropriate model for the *auto-correlation structure*. Linear relationships, so this amounts to picking the appropriate **lag** or **order** of the model

- Should current Y be predicted by Y an hour ago, or should we also include the effect of Y two hours ago? Three hours ago? 24 hours ago?

Y

y_1

y_2

y_3

y_4

y_5

y_6

y_7

y_8

\dots

y_T

Lags

Y	Y at lag 1
y_1	
y_2	y_1
y_3	y_2
y_4	y_3
y_5	y_4
y_6	y_5
y_7	y_6
y_8	y_7
...	...
y_T	y_{T-1}
	y_T

Lags

Y	Y at lag 1	Y at lag 2
y_1		
y_2	y_1	
y_3	y_2	y_1
y_4	y_3	y_2
y_5	y_4	y_3
y_6	y_5	y_4
y_7	y_6	y_5
y_8	y_7	y_6
...
y_T	y_{T-1}	y_{T-2}
	y_T	y_{T-1}
		y_T

How to choose an appropriate Time-Series Model?

Box-Jenkins Method (1970). Aim: Obtain a model with **white noise** residuals (error term contains no information about the future)

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- **Autocorrelation function (ACF).**
Raw/marginal correlation between lagged versions of our variable y_t and y_{t-k}

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Diagnostic Tools:

- **Autocorrelation function (ACF)**. Raw/marginal correlation between lagged versions of our variable y_t and y_{t-k}
- **Partial autocorrelation functions (PACF)**. Partial correlation between y_t and y_{t-k} when controlling for the effect of intermediate observations y_{t-1} to y_{t-k+1} .

ACF and PACF explained

ACF Lag-1

$cor(Y_t, Y_{t-1})$

Y	Y at lag 1	Y at lag 2
y_1		
y_2	y_1	
y_3	y_2	y_1
y_4	y_3	y_2
y_5	y_4	y_3
y_6	y_5	y_4
y_7	y_6	y_5
y_8	y_7	y_6
...
y_T	y_{T-1}	y_{T-2}
	y_T	y_{T-1}
		y_T

ACF and PACF explained

ACF Lag-2

$cor(Y_t, Y_{t-2})$

Y	Y at lag 1	Y at lag 2
y_1		
y_2	y_1	
y_3	y_2	y_1
y_4	y_3	y_2
y_5	y_4	y_3
y_6	y_5	y_4
y_7	y_6	y_5
y_8	y_7	y_6
...
y_T	y_{T-1}	y_{T-2}
	y_T	y_{T-1}
		y_T

ACF and PACF explained

PACF Lag-2

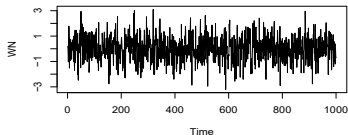
$$\text{cor}(Y_t, Y_{t-2} \mid Y_{t-1})$$

Y	Y at lag 1	Y at lag 2
y ₁		
y ₂	y ₁	
y ₃	y ₂	y ₁
y ₄	y ₃	y ₂
y ₅	y ₄	y ₃
y ₆	y ₅	y ₄
y ₇	y ₆	y ₅
y ₈	y ₇	y ₆
...
y _T	y _{T-1}	y _{T-2}
	y _T	y _{T-1}
		y _T

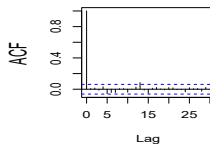
Similar to a regression coefficient – controlling for Y_{t-1}

Sequence, ACF and PACF

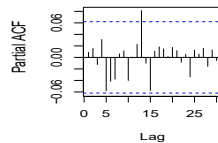
White Noise process



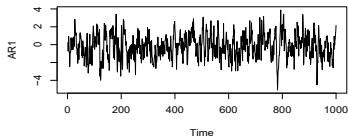
Series WN



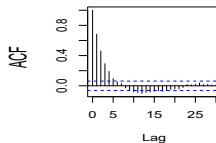
Series WN



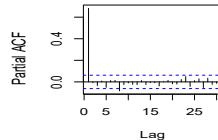
First-order AR process



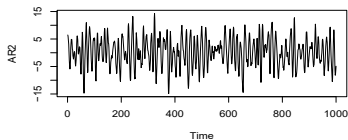
Series AR1



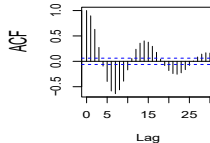
Series AR1



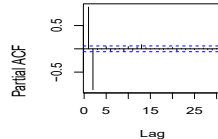
Second-order AR process



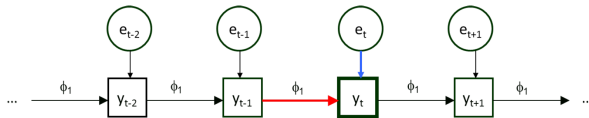
Series AR2



Series AR2

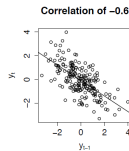
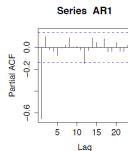
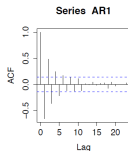
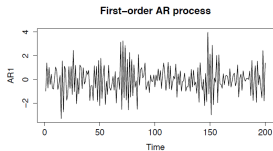
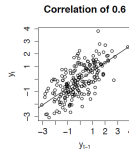
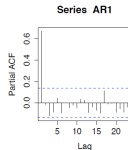
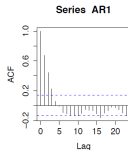
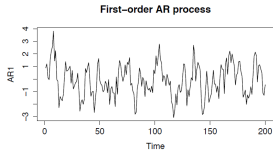


AR(1) Model

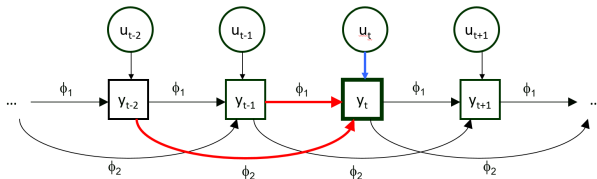


$$y_t = \phi_1 y_{t-1} + e_t$$

$$\phi_1 = \{0.6; -0.6\}$$

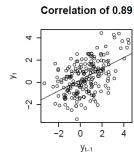
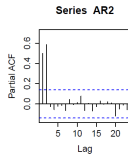
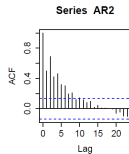
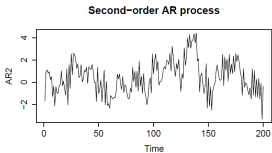


AR(2) Model



$$y_t = \phi_1 y_{t-1} + \phi_2 y_{t-2} + e_t$$

$$\phi_1 = 0.2; \phi_2 = 0.6$$



General approach:

- 1 Check for trends and seasonal components. Remove them if present.
- 2 Plot ACF and PACF. Choose an appropriate autocorrelation structure. Obtain residuals
- 3 Plot ACF and PACF of residuals. Check if white noise. If not, repeat step 2.

General approach:

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Sidenote: Sometimes referred to as AR(I)MA modeling. We focus here on autoregressive (AR) terms, but moving average (MA) terms also possible

MA(1) model:

$$y_t = c + \varepsilon_t + \theta \varepsilon_{t-1}$$

Granger and Morris (1976): An AR process is a **momentum effect in a random variable that varies smoothly over time**

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We think about a variable with a stable **equilibrium position**. The errors represent “shocks” that push the system away from equilibrium. The AR parameter describes the “carry-over” or “regulation” of those shocks from one moment to the next.

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AR(1) model: The **auto-regressive parameter** pulls the system **back** to equilibrium. The bigger the **absolute value**, the slower the system is to return to baseline

Impulse Response Functions help us understand qualitative behaviour

- 1 Pick an interesting value for Y_0
- 2 Take your model + parameters
- 3 Calculate expected value of Y_1
- 4 Repeat and plot to visualize time-evolution of the system

A simple discrete-time model

Impulse Response Functions help us understand qualitative behaviour

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AR(1) Model

$$E[Y_1] = \phi Y_0$$

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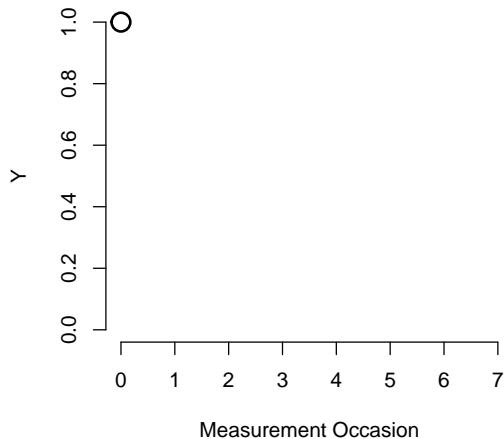
AR(1) Model

$$E[Y_1] = \phi Y_0$$

with

- $\phi = .5$
- $Y_0 = 1$

Impulse Response Function



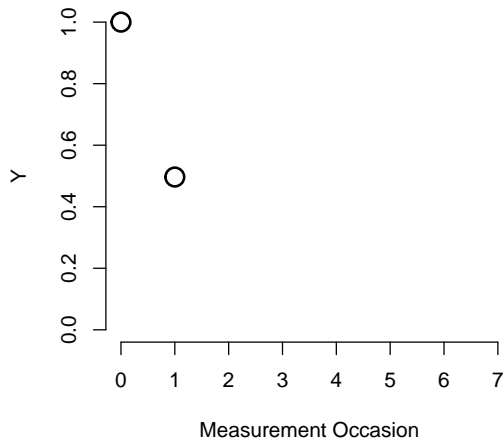
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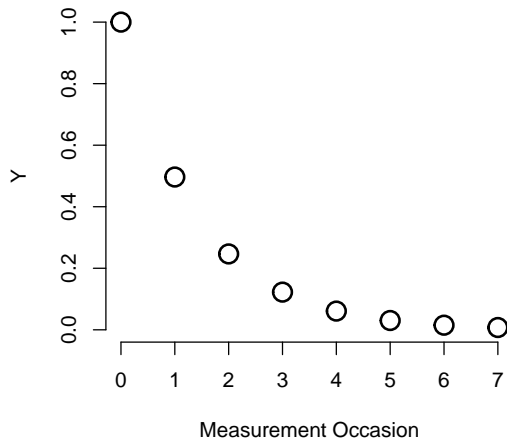
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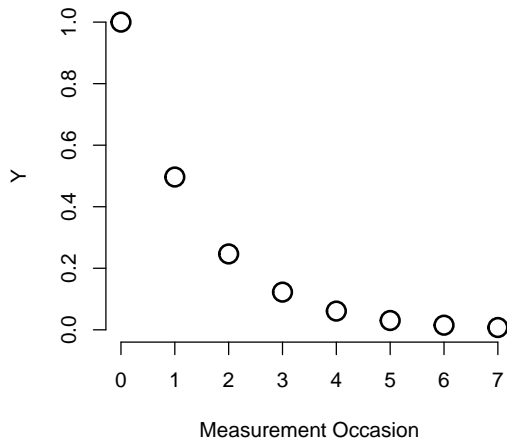
AR(1) Model

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Impulse Response Function



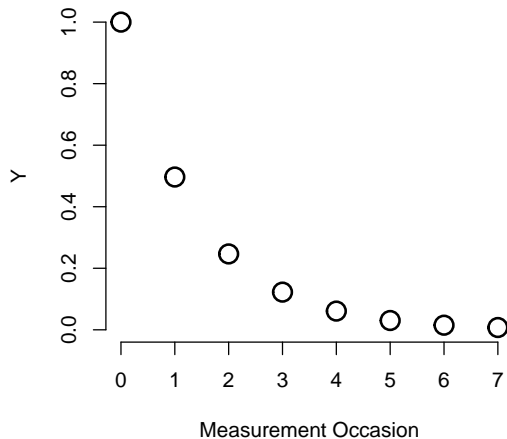
AR(1) Model

$$E[Y_2] = \phi \times \phi Y_0$$

with

- $\phi = .5$
- $Y_0 = 1$

Impulse Response Function



AR(1) Model

$$E[Y_\tau] = \phi^\tau Y_0$$

with

- $\phi = .5$
- $Y_0 = 1$

Time-series analysis focuses on modeling *stationary* processes.

Stationary processes are those for which the mean, variance, and auto-correlation structure **stay the same over our window of observation**.

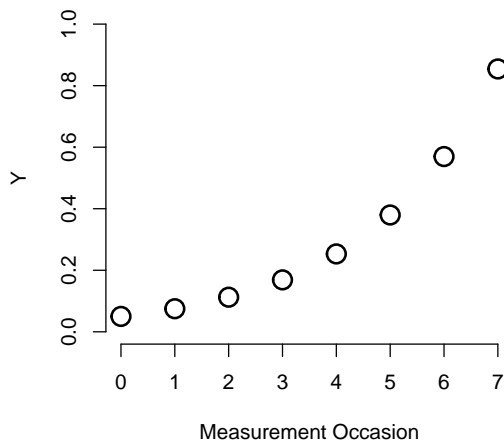
- Trends can produce non stationarity (changes in mean over time), so have to be removed
 - Failing to remove a trend if present produces bias... but more on that later!
- Lagged regression parameters stay fixed across waves
 - Always necessary in some form - otherwise impossible to estimate!
- Only certain parameter values allowed
 - For AR(1) $-1 < \phi < 1$

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 - Always necessary in some form - otherwise impossible to estimate!
- Only certain parameter values allowed
 - For AR(1) $-1 < \phi < 1$
 - AR(1) interpretation: system that gets pushed away from equilibrium by shocks but always returns to equilibrium

Impulse Response Function: Non-Stationary



AR(1) Model

$$E[Y_\tau] = \phi^\tau Y_0$$

with

- $\phi = 1.5$
- $Y_0 = 0.05$

Summary: Univariate Auto-regressive models

After removing of modeling trends and stationarity, we can fit an AR(p) model

$$Y_t = c + \phi_1 Y_{t-1} + \cdots + \phi_p Y_{t-p} + \varepsilon_t$$

where:

- c is the intercept term
- ϕ_p is the auto-regressive effect of order p
- ε_t is a white-noise Gaussian error term $\varepsilon_t \sim N(0, \sigma_\varepsilon)$
- We try to choose an appropriate p using the ACF and PACF

A stationary AR(p) process is one which fluctuates around a *stable equilibrium* i.e. long-run mean value μ

- For an AR(1), the mean is given by $\mu = \frac{c}{1-\phi}$

- Time series analysis: Univariate
- **Time series analysis: Multivariate**
- Multilevel time series analysis
- Extensions and Advanced Issues
- Discussion

Typically we have ILD that consist of **more than one** variable:

- mother's depression and child's disruptive behaviour
- stress and anxiety
- physical activity and happiness
- sleep quality and mood
- ...

In that situation, we might be interested in what time-series researchers call “**lead-lag relations**” between these variables

To analyze these relationships, we can use the multivariate generalization of the AR(p)

Known as the **Vector Autoregressive Model** or VAR(p)

$$\mathbf{Y}_t = \mathbf{c} + \Phi_1 \mathbf{Y}_{t-1} + \cdots + \Phi_p \mathbf{Y}_{t-p} + \epsilon_t$$

where:

- \mathbf{c} is a $q \times 1$ vector of intercepts
- Φ_p is a $q \times q$ matrix of auto-regressive and cross-lagged effects
- ϵ_t is a vector of multivariate Gaussian errors $\epsilon_t \sim N(0, \Sigma_\epsilon)$

A simple multivariate time-series model

The VAR(1) model is a multivariate version of the popular AR(1) model. We can write it as a set of equations:

$$A_t = c_A + \phi_{11}A_{t-1} + \phi_{12}S_{t-1} + e_{A_t}$$

$$S_t = c_S + \phi_{21}A_{t-1} + \phi_{22}S_{t-1} + e_{S_t}$$

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or in **matrix form**

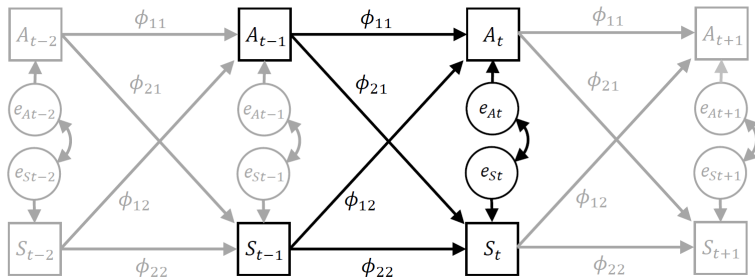
$$\begin{bmatrix} A_t \\ S_t \end{bmatrix} = \begin{bmatrix} c_A \\ c_S \end{bmatrix} + \begin{bmatrix} \phi_{11} & \phi_{12} \\ \phi_{21} & \phi_{22} \end{bmatrix} \begin{bmatrix} A_{t-1} \\ S_{t-1} \end{bmatrix} + \begin{bmatrix} e_{A_t} \\ e_{S_t} \end{bmatrix}$$

where:

ϕ_{11} and ϕ_{22} are the **autoregressive coefficients**

ϕ_{12} and ϕ_{21} are the **cross-lagged coefficients**

Bivariate VAR(1)



Researchers often interpret the cross-lagged parameters in terms of **causal dominance** or **Granger causal** relationships. Note: the **standardized cross-lagged regression coefficients** should be used!

VAR(1) models very popular: easy to fit and visualize the output

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Become an increasingly popular tool since the popularity of the **network theory** of psychopathology (Borsboom, 2017)

- Idea: Mental disorders arise due to direct causal connections between themselves, not due to some latent "disease" variable
- Typical Empirical Application: Attempt to learn about network structure using statistical models which encode direct conditional dependencies

VAR(1) models very popular: easy to fit and visualize the output

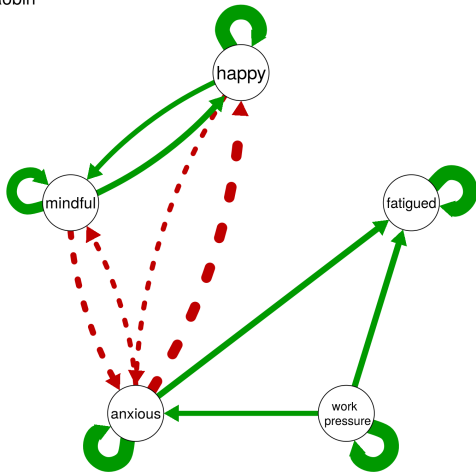
Become an increasingly popular tool since the popularity of the **network theory** of psychopathology (Borsboom, 2017)

- Idea: Mental disorders arise due to direct causal connections between themselves, not due to some latent "disease" variable
- Typical Empirical Application: Attempt to learn about network structure using statistical models which encode direct conditional dependencies
- VAR(1) gives the simplest possible directed "network structure" from time-series data

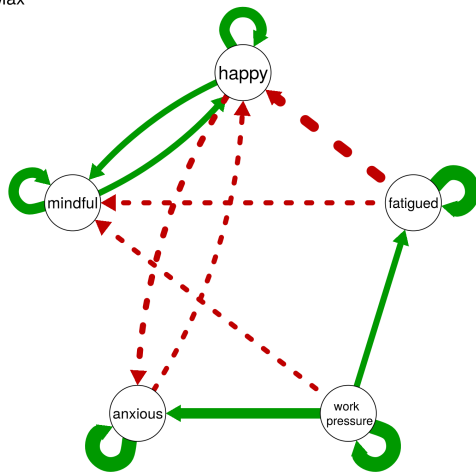
General approach popular in other fields, special case of "Dynamic Bayesian Networks"

VAR(1) networks

Robin



Max

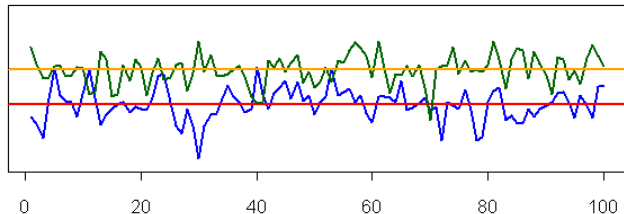


VAR(1) networks

Almost always a "dynamical network" in psychology is just a VAR(1) model with it's lagged parameters plotted

Remember: Qualitative dynamics same as the AR(1)

- Single resting state, which we either always return to or always explode away from



- Time series analysis: Univariate
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Single-subject time series data allows us to study (for a *stationary process*):

- Lagged relationships between a variable and itself: **autoregression**
- Lagged relationships between different variables: **cross-lagged relationships**

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- individual differences in **autoregression**
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- individual differences in **autoregression**
- individual differences in **cross-lagged relationships**

If we use multilevel modeling for this, we could refer to it as **multilevel time series analysis**, or **dynamic multilevel modeling**.

Creating lagged predictors

ID

1

1

1

1

1

2

2

2

2

2

...

N

N

N

Creating lagged predictors

ID	y_{it}
1	y_{11}
1	y_{12}
1	y_{13}
1	...
1	y_{1T}
2	y_{21}
2	y_{22}
2	y_{23}
2	...
2	y_{2T}
...	...
N	y_{N1}
N	y_{N2}
N	y_{N3}

Creating lagged predictors

ID	y_{it}	y_{it-1}
1	y_{11}	
1	y_{12}	y_{11}
1	y_{13}	y_{12}
1
1	y_{1T}	y_{1T-1}
2	y_{21}	
2	y_{22}	y_{21}
2	y_{23}	y_{22}
2
2	y_{2T}	y_{2T-1}
...
N	y_{N1}	
N	y_{N2}	y_{N1}
N	y_{N3}	y_{N2}

Creating lagged predictors

ID	y_{it}	y_{it-1}	x_{it-1}
1	y_{11}		
1	y_{12}	y_{11}	x_{11}
1	y_{13}	y_{12}	x_{12}
1
1	y_{1T}	y_{1T-1}	x_{1T-1}
2	y_{21}		
2	y_{22}	y_{21}	x_{21}
2	y_{23}	y_{22}	x_{22}
2
2	y_{2T}	y_{2T-1}	x_{2T-1}
...
N	y_{N1}		
N	y_{N2}	y_{N1}	x_{N1}
N	y_{N3}	y_{N2}	x_{N2}

Level 1 model:

$$NA_{it} = c_i + \phi_i NA_{i,t-1} + \zeta_{it}$$

Inertia research based on multilevel AR(1) models

Level 1 model:

$$NA_{it} = c_i + \phi_i NA_{i,t-1} + \zeta_{it}$$

Level 2 model:

$$c_i = \gamma_{00} + u_{0i}$$

$$\phi_i = \gamma_{01} + u_{1i}$$

Inertia research based on multilevel AR(1) models

Level 1 model:

$$NA_{it} = c_i + \phi_i NA_{i,t-1} + \zeta_{it}$$

Level 2 model:

$$c_i = \gamma_{00} + u_{0i}$$

$$\phi_i = \gamma_{01} + u_{1i}$$

This research line was initiated by **Suls, Green and Hillis (1998)**, and continued by the group of **Kuppens**.

The focus is on individual differences in the **autoregressive parameter** ϕ_i (=inertia, carry-over, regulatory weakness), which is shown to be:

- positively related to current depression, neuroticism, and being female
- predictive of later depression (Kuppens and Koval)

Dynamic networks based on multilevel VAR(1) models

Level 1 model:

$$y_{1it} = c_{1i} + \phi_{11i}y_{1it-1} + \cdots + \phi_{1ki}y_{kit-1} + \zeta_{1it}$$

$$y_{2it} = c_{2i} + \phi_{21i}y_{1it-1} + \cdots + \phi_{2ki}y_{kit-1} + \zeta_{2it}$$

...

$$y_{kit} = c_{ki} + \phi_{k1i}y_{1it-1} + \cdots + \phi_{kki}y_{kit-1} + \zeta_{kit}$$

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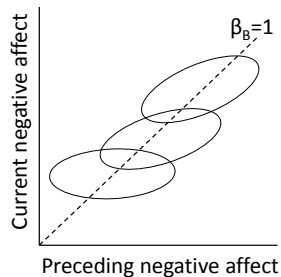
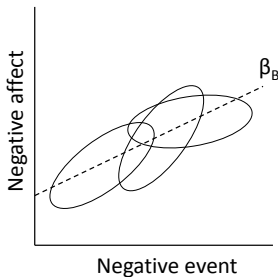
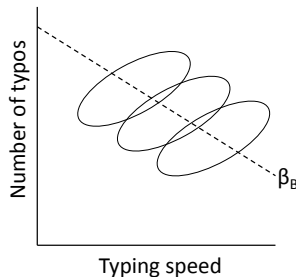
Initiated by **Bringmann et al. (2013)**, and further popularized by the software from **Sacha Epskamp**.

The focus is on **cross-lagged parameters** between variables (=nodes; typically symptoms), and on measures based on these (e.g., centrality).

Main idea is that **stronger connections** lead to an **increased risk** of developing and maintaining psychopathology.

Between-person differences in within-person slopes

- Studying a collection of within-person processes
- We want to know the average within-person process (fixed effects) and how these processes differ between people (random parts)
- "Decomposition" of data into within- and between- person variance components



Taken from Hamaker and Grasman (2014).

Decomposition into a between part and a within part

$$PA_{it} = \mu_{PA,i} + PA_{it}^*$$

$$NA_{it} = \mu_{NA,i} + NA_{it}^*$$

Decomposition

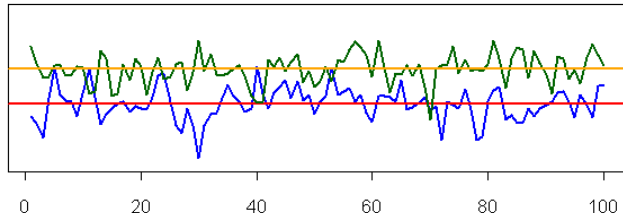
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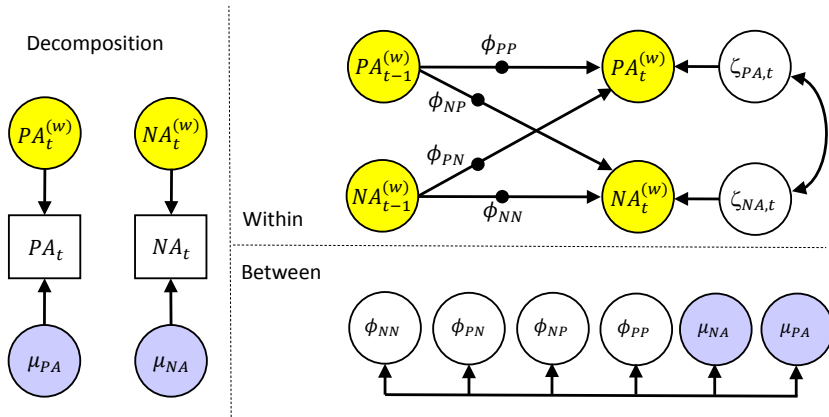
$$NA_{it} = \mu_{NA,i} + NA_{it}^*$$

where

- $\mu_{PA,i}$ and $\mu_{NA,i}$ are the individual's **means** on PA and NA (i.e., baseline, trait, or equilibrium scores) \Rightarrow between-person part
- PA_{it}^* and NA_{it}^* are the **within-person centered** (cluster-mean centered) scores \Rightarrow within-person part



Bivariate model: Multilevel VAR(1) model



Within-person level model

Lagged within-person model:

$$\begin{aligned}PA_{it}^* &= \phi_{PP,i}PA_{i,t-1}^* + \phi_{PN,i}NA_{i,t-1}^* + \zeta_{PA,it} \\ NA_{it}^* &= \phi_{NN,i}NA_{i,t-1}^* + \phi_{NP,i}PA_{i,t-1}^* + \zeta_{NA,it}\end{aligned}$$

where

- $\phi_{PP,i}$ is the **autoregressive parameter** for PA (i.e., inertia, carry-over)
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- $\zeta_{PA,it}$ is the **innovation** for PA (residual, disturbance, dynamic error)
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Within-person level model

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- $\zeta_{NA,it}$ is the **innovation** for NA (residual, disturbance, dynamic error)

Residual variances and covariances:

$$\begin{bmatrix} \zeta_{PA,it} \\ \zeta_{NA,it} \end{bmatrix} \sim MN \left[\begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \theta_{11} & \\ \theta_{21} & \theta_{22} \end{bmatrix} \right]$$

Between-person level model

Between level: fixed and random effects

$$\begin{bmatrix} \mu_{PA,i} \\ \mu_{NA,i} \\ \phi_{PP,i} \\ \phi_{PN,i} \\ \phi_{NP,i} \\ \phi_{NN,i} \end{bmatrix} = \begin{bmatrix} \gamma_P \\ \gamma_N \\ \gamma_{PP} \\ \gamma_{PN} \\ \gamma_{NP} \\ \gamma_{NN} \end{bmatrix} + \begin{bmatrix} u_{P,i} \\ u_{N,i} \\ u_{PP,i} \\ u_{PN,i} \\ u_{NP,i} \\ u_{NN,i} \end{bmatrix} \quad \mathbf{u}_i \sim MN(\mathbf{0}, \Psi)$$

Where:

- γ_P to $\gamma_{NN} \Rightarrow$ fixed effects
- $u_{P,i}$ to $u_{NN,i} \Rightarrow$ random effects

Between-person level model

Between level: fixed and random effects

$$\begin{bmatrix} \mu_{PA,i} \\ \mu_{NA,i} \\ \phi_{PP,i} \\ \phi_{PN,i} \\ \phi_{NP,i} \\ \phi_{NN,i} \end{bmatrix} = \begin{bmatrix} \gamma_P \\ \gamma_N \\ \gamma_{PP} \\ \gamma_{PN} \\ \gamma_{NP} \\ \gamma_{NN} \end{bmatrix} + \begin{bmatrix} u_{P,i} \\ u_{N,i} \\ u_{PP,i} \\ u_{PN,i} \\ u_{NP,i} \\ u_{NN,i} \end{bmatrix} \quad \mathbf{u}_i \sim MN(\mathbf{0}, \Psi)$$

Where:

- γ_P to $\gamma_{NN} \Rightarrow$ fixed effects
- $u_{P,i}$ to $u_{NN,i} \Rightarrow$ random effects

Parameters estimated at this level are:

- 6 fixed effects (i.e., γ 's)
- 6 variances for random effects (i.e., diagonal elements of Ψ)
- 15 covariances between the random effects (i.e., off-diagonal elements in Ψ)

Why Bayesian Estimation?

Bayesian approaches to fitting multilevel time-series models have become increasingly popular, for a number of reasons:

- In principle can handle missing observations in a tidier way
- Simultaneous estimation of entire model, rather than variable-by-variable approach
- Can incorporate measurement models / latent variables
- Can more easily output within-person standardized cross-lagged parameters
- Bayesian approaches can be implemented in BUGS/JAGS, STAN, and have been implemented in DSEM in Mplus

Intermezzo on Bayesian analysis

Bayesian analysis is based on combining the **density of the data** with a **prior distribution** for the unknown parameters, to get a **posterior distribution** of these parameters.

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Posterior distribution of θ

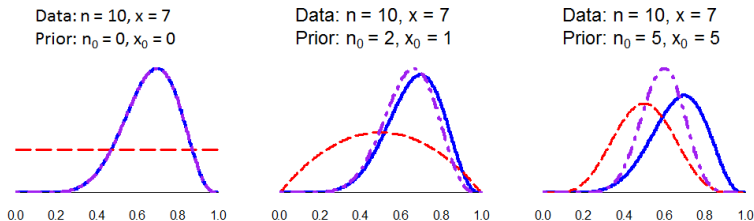
$$p(\theta|y) = \frac{f(y|\theta)p(\theta)}{f(y)}$$

where:

- $f(y|\theta)$ be the **density of the data** y given the parameters θ (also referred to as the likelihood)
- $p(\theta)$ be the **prior distribution** of the parameter(s) θ , which the user needs to specify
- $\int f(y, \theta)d\theta = f(y)$ is the **marginal density**, which can be ignored (because it is a constant)

Intermezzo on Bayesian analysis

Density (blue), prior (red), and posterior (purple):



When the prior is flat (no information), the posterior is identical to the likelihood.

If you have prior knowledge, you can add this to the equation by specifying a prior that reflects this.

For each to be estimated parameter, a prior needs to be specified. In the lab we'll aim to specify uninformative priors.

Intermezzo on Bayesian analysis: Convergence

Bayesian analysis is (often) based on using an **MCMC algorithm** which iteratively **samples** the parameters from their conditional posteriors.

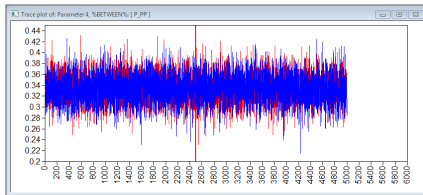
We have to check whether the analysis has **converged** (or: whether there are signs it did **not** converge).

Tools we use for this are:

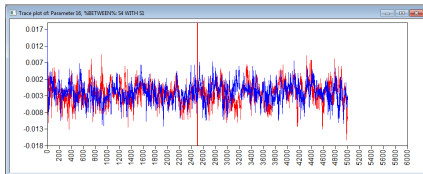
- Multiple chains; multiple runs of the analysis with different starting values.
- These chains should end up at approximately the same estimates.
- Burnin: Part of the iterations (before convergence) are discarded, leaving only 'converged' samples.
- Plots of the chains (fat hairy caterpillars), density plots (should look smooth and normal-ish), gelman rubin statistic: should be very close to 1.

Intermezzo on Bayesian analysis: Trace plots

This looks good (lazy, fat caterpillar):



This looks less good but not really bad; just needs more samples:



	N=1	multilevel
uni-variate	<ul style="list-style-type: none">- arima in R- State Space Modeling software- Openmx- Bayesian modeling software	<ul style="list-style-type: none">- any multilevel software- MLvar package in R- Bayesian modeling software
some-what multi-variate	<ul style="list-style-type: none">- VARS package in R- State Space Modeling Software- Openmx- Bayesian modeling software	<ul style="list-style-type: none">- any multilevel software- MLVar package in R- Bayesian modeling software
multi-variate	<ul style="list-style-type: none">- State Space Modeling Software (mkfm6; Ox; fkf, dlm, KFAS, and MARSS in R)- Bayesian software (Winbugs, Openbugs, JAGS, STAN)	<ul style="list-style-type: none">- Bayesian software (Winbugs, Openbugs, JAGS, STAN)

Advantages of Multi-Level Time Series Models:

- Allow to model within-person and between-person structure simultaneously
- Regularizes estimates of within-person parameters
 - Person-specific parameters get “pulled” towards the mean. Estimates have lower variance across samples
- The more similar people are, and the more people we have, the less observations we need per person

Multilevel vs Single-Subject Time Series

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Disadvantages of Multi-Level Time Series Models:

- Need to assume the same type of time series model (or a very general model) for everyone
- Regularizes estimates of within-person parameters
 - If we have enough time points, and we really care about the individual-specific parameters, we don't want this to happen
 - If people are not that similar, we might get a very strange/uninformative “average” model

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Comparing cross-lagged parameters

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Schuurman, Ferrer, Boer-Sonnenschein and Hamaker (2016) discuss four forms of **standardization in multilevel models**, using:

- total variance (i.e., grand standardization)
- between-person variance (i.e., between standardization)
- average within-person variance
- within-person variance (i.e., within standardization)

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Conclusion: last form is most meaningful, as it **parallels standardizing when $N=1$** .

Standardized fixed effect should be the **average standardized within-person effect**.

Within the model, variables are **within-person centered**

Decomposition into a between part and a within part

$$PA_{it} = \mu_{PA,i} + PA_{it}^*$$

$$NA_{it} = \mu_{NA,i} + NA_{it}^*$$

This also makes it easier to obtain within-person standardized lagged effects.

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This also makes it easier to obtain within-person standardized lagged effects.

If we have a trend, i.e., some linear or quadratic function of time t , then we typically want to **de-trend per person**.

Be careful: not always implemented in multilevel software, and if you de-trend before fitting the model, the means for every person will be zero!

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Deterministic Trend: variables are a direct function of time, which we treat as an exogenous variable

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Solution: fit a linear a trend to "de-trend"

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$$y_t = c + \phi y_{t-1} + bT + e_t$$

Solution: fit a linear a trend to "de-trend"

Stochastic trend: non-stationarity is produced by the dynamics of the endogenous variable

$$y_t = c + 1.5y_{t-1} + e_t$$

There are actually two distinct mechanisms which can produce **non-stationarity**

Deterministic Trend: variables are a direct function of time, which we treat as an exogenous variable

$$y_t = c + \phi y_{t-1} + bT + e_t$$

Solution: fit a linear a trend to "de-trend"

Stochastic trend: non-stationarity is produced by the dynamics of the endogenous variable

$$y_t = c + 1.5y_{t-1} + e_t$$

Solution: *differencing*. $y^* = y_t - y_{t-1}$, then y^* is stationary.

Failing to de-trend when there's a **deterministic trend** present results in bias

But **erroneously de-trending** when actually there is a **stochastic trend** can also lead to bias!

You can test for the presence of a stochastic trend using the *Augmented Dickey-Fuller* test. More details in Ryan & Haslbeck (in preparation)

Time-Varying VAR(1) models:

- Haslbeck, Bringmann & Waldorp (2020)
- Assumes “local stationarity” - all parameters can be smooth functions of time
- Mostly single-subject, needs even more time-points

Regime-Switching Models:

- Time-series models where the system “switches” between different states with different parameters
- (Hidden) Markov Models, Threshold VAR models
- See Hamaker, Grasman, Kamphuis (2010); Haslbeck* & Ryan* (2021); dynr (Ou, Hunter, Chow, 2017)

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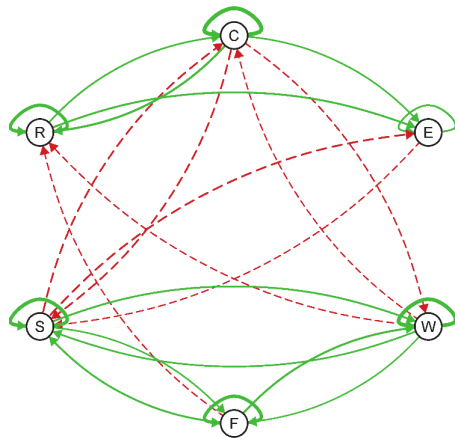
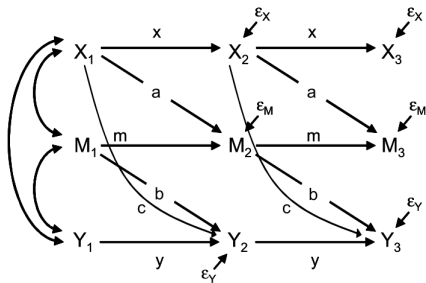
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Multilevel Bayesian setting: things are trickier. **DIC** is the Bayesian information criteria, but model comparison can be tricky due to irregularities in how model complexity is counted and stability of DIC across samples.

Very much a work in progress. Personal tip: You can always use out-of-sample prediction error to compare models!

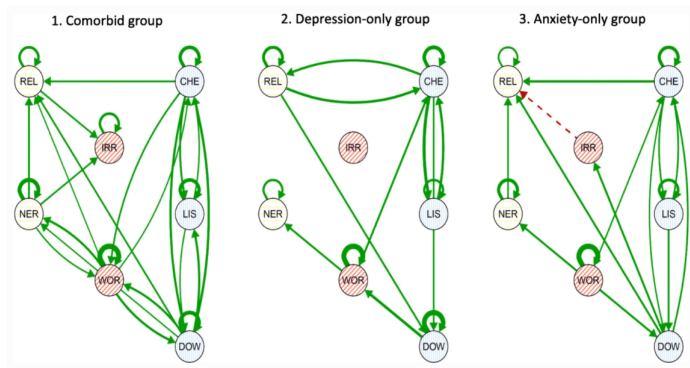
Level-1 Mediation and Network Analysis

We may also be interested in *direct*, *indirect* and *total effects across lags*. Often the interest in *psychological network analysis*



Level-1 Mediation and Network Analysis

Groen, Ryan, Wigman et al (2020): Average within-person VAR(1) network structure of different “symptom-state” items in ESM data, across three groups: Comorbid, Anxiety-only and Depression-only.



No evidence that the comorbid group displayed higher indirect effects through “bridge symptoms” than the other groups. All analyses in DSEM

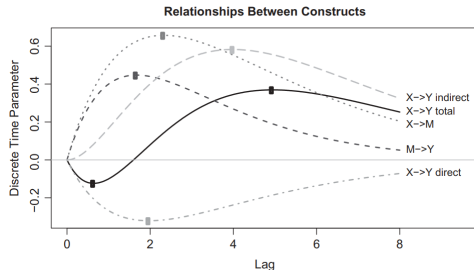
The Time-Interval Problem

VAR(1) models are **discrete time**: Do not account in any way for time-interval information

All lagged regression models (potentially) suffer from the time-interval problem.

Effects can change sign, size and relative ordering depending on how measurements are spaced in time (Kuiper & Ryan, 2018)

More on this **tomorrow!**



- Time series analysis
- Multilevel time series analysis
- DSEM application: Multilevel VAR(1) model
- Extensions and Issues
- **Discussion**

Time-series analysis aims to capture the structure of within-person variation over time when we have many repeated measures

When we have ILD from multiple individuals, **multi-level time series models** can be used to try and explore how within-person structure varies across people

The VAR(1) is just one type of time-series model which happens to be very popular - simple, lag-1 linear relationships.

The qualitative behaviour of this system is the same as the AR(1), but multivariate: fluctuating around a single equilibrium position.

Of course, big issue is: which time-series models are substantively interesting?

- Which captures/reproduces patterns that are interesting?
- How can this be driven by theory?

Approach direct (and causal) interpretation with care

Tip: Pick a model and simulate time-series data from it (e.g. in R) - get a feel for what types of patterns it produces

These models are just tools. Proper use must be informed by theory (Haslbeck*, Ryan*, Robinaugh* et al, in press)

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