

Development / longitudinal / repeated measures / time series / (not cross-sectional)

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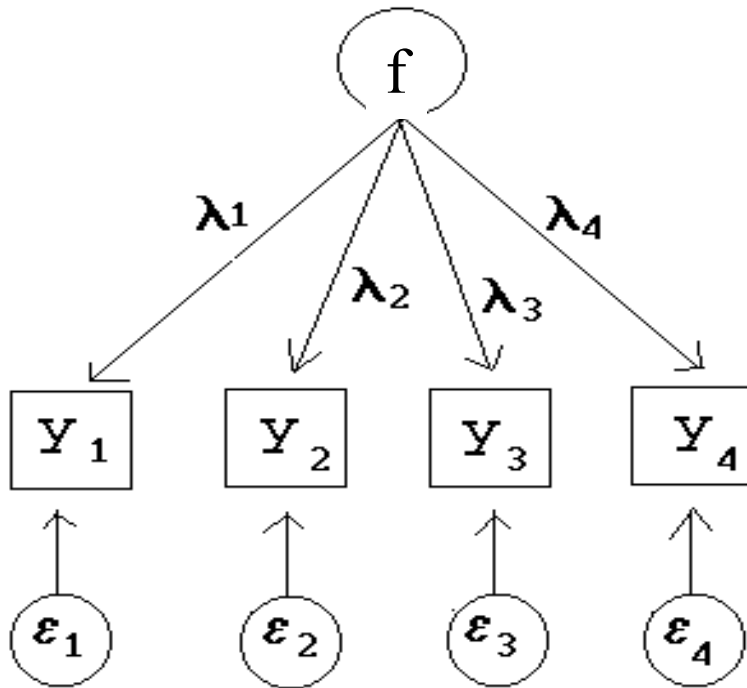
Netherlands Twin Register

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General multivariate model for 1 person: Regression of observed variables (y) on latent variables (f)



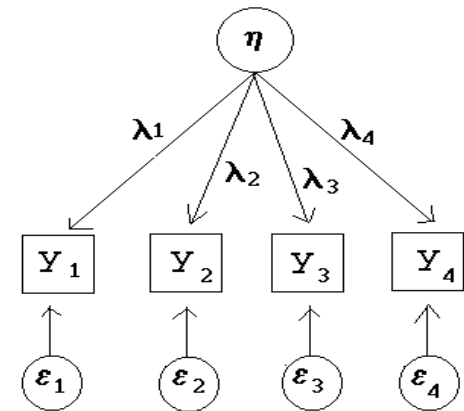
Covariance among variables is accounted for by a smaller number of latent factors.

Model: $y = \Lambda f + e$,

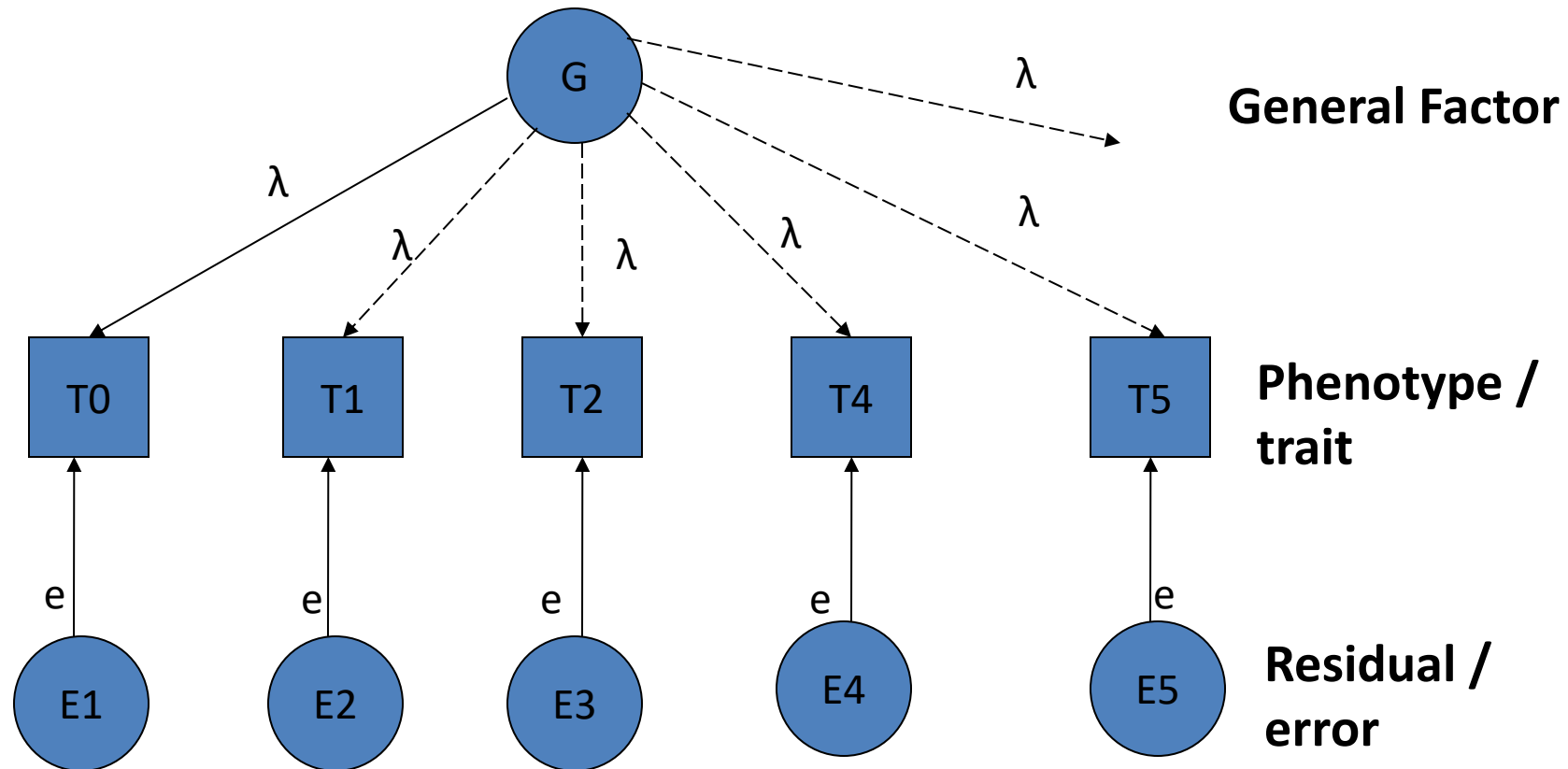
where y = observed variables
 f = (latent) factor score(s)
 e = unique factor / error
 Λ = matrix of factor loadings

Structural equation models.

Sometimes $x = \Lambda f + e$ is referred to as the measurement model, and relations among latent factors as the structural equation model.

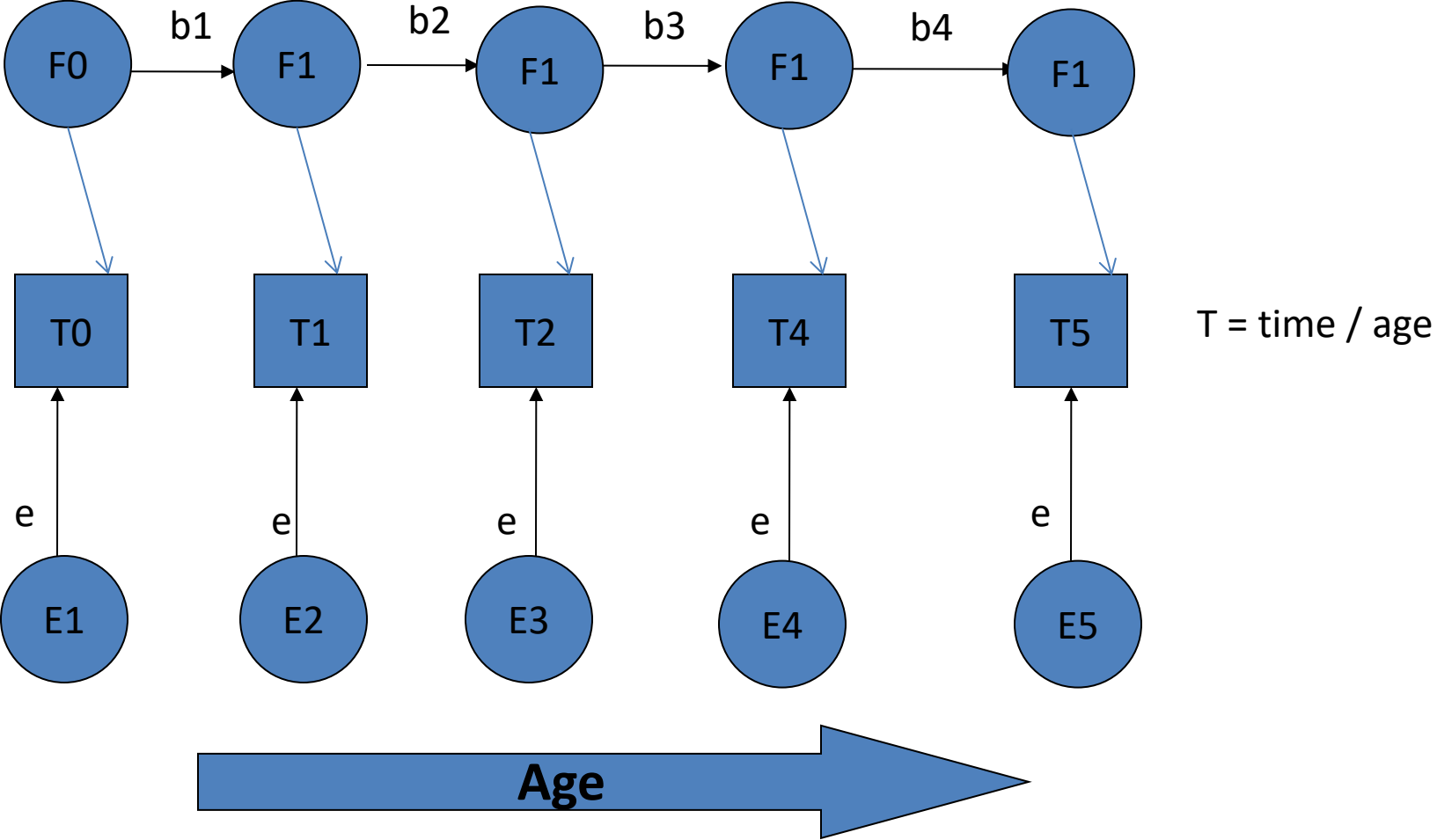


Multivariate model



If $T = \text{time}$ and N of time points get large, what will happen?

Developmental model



Two general approaches to longitudinal modeling (not mutually exclusive)

Markov models:

(Vector) autoregressive models for continuous data

(Hidden) Markov transition models discrete data

Growth curve models:


Focus on linear and non-linear growth curves

Typically multilevel or random effects model

Which to use?

Use the model that fit the theory & hypotheses

Growth curve modeling ? If you're interested in growth trajectories. Linear or non-linear:

Twin Research (2000) 3, 165-177
© 2000 Macmillan Publishers Ltd All rights reserved 1369-0523/00 \$15.00 
www.nature.com/tr

Structured latent growth curves for twin data

Michael C Neale¹ and John J McArdle²

Autoregressive modeling ? If you're interested in e.g. the genetic stability of a trait (e.g combining data).

Psychological Medicine (2015), 45, 1039-1049. © Cambridge University Press 2014
doi:10.1017/S003329171400213X

ORIGINAL ARTICLE

Stability in symptoms of anxiety and depression as a function of genotype and environment: a longitudinal twin study from ages 3 to 63 years

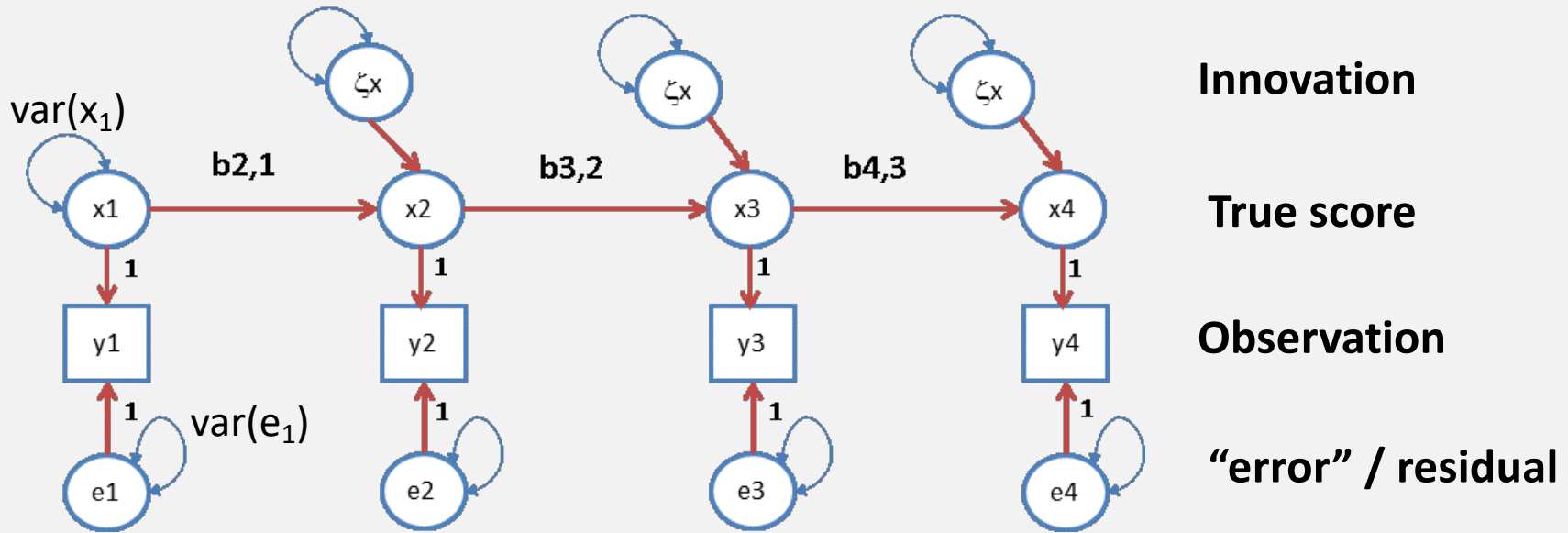
M. G. Nivard^{1,2*}, C. V. Dolan^{1,3}, K. S. Kendler⁴, K.-J. Kan¹, G. Willemsen^{1,5},
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M. Bartels^{1,5}, C. M. Middeldorp^{1,2,7†} and D. I. Boomsma^{1,2,5†}

Do the Genetic or Environmental Determinants of Anxiety and Depression Change with Age? A Longitudinal Study of Australian Twins

Nathan A. Gillespie¹, Katherine M. Kirk¹, David M. Evans¹, Andrew C. Heath², Ian B. Hickie³, and Nicholas G. Martin¹

Can be combined (Nathan Gillespie)

First order autoregression model.



Innovation

True score

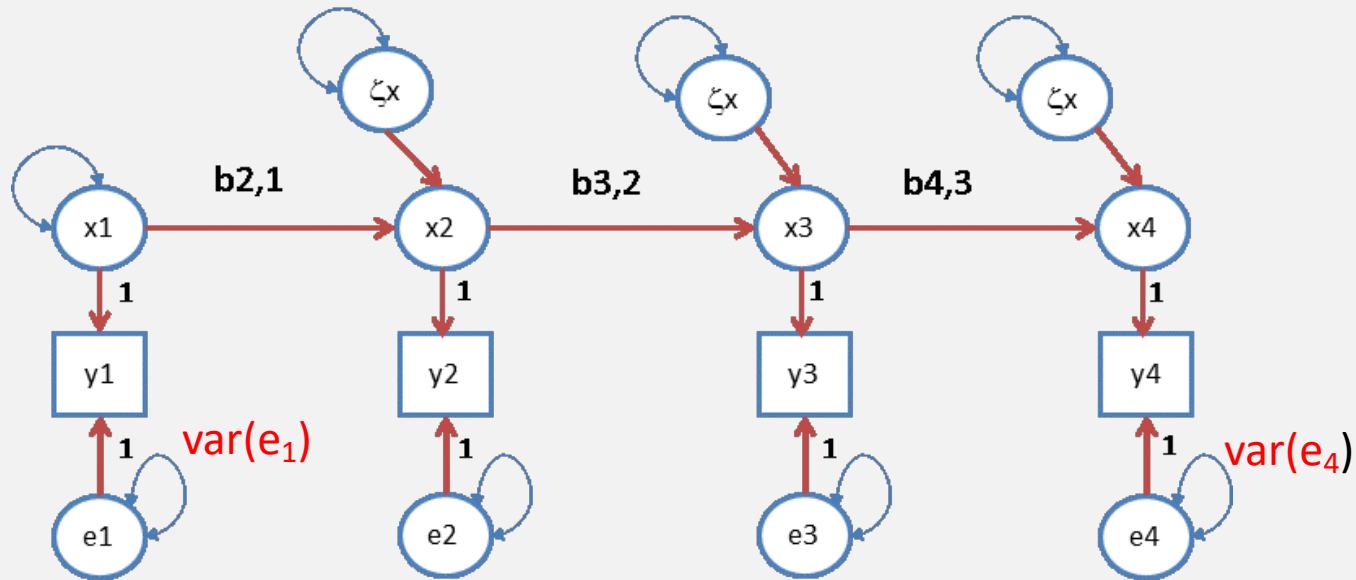
Observation

“error” / residual

$$y_{ti} = b_{0t} + x_{ti} + e_{ti} \quad (\text{measurement model})$$

$$x_{ti} = b_{t-1,t} x_{t-1i} + \zeta x_{ti} \quad (\text{structural model})$$

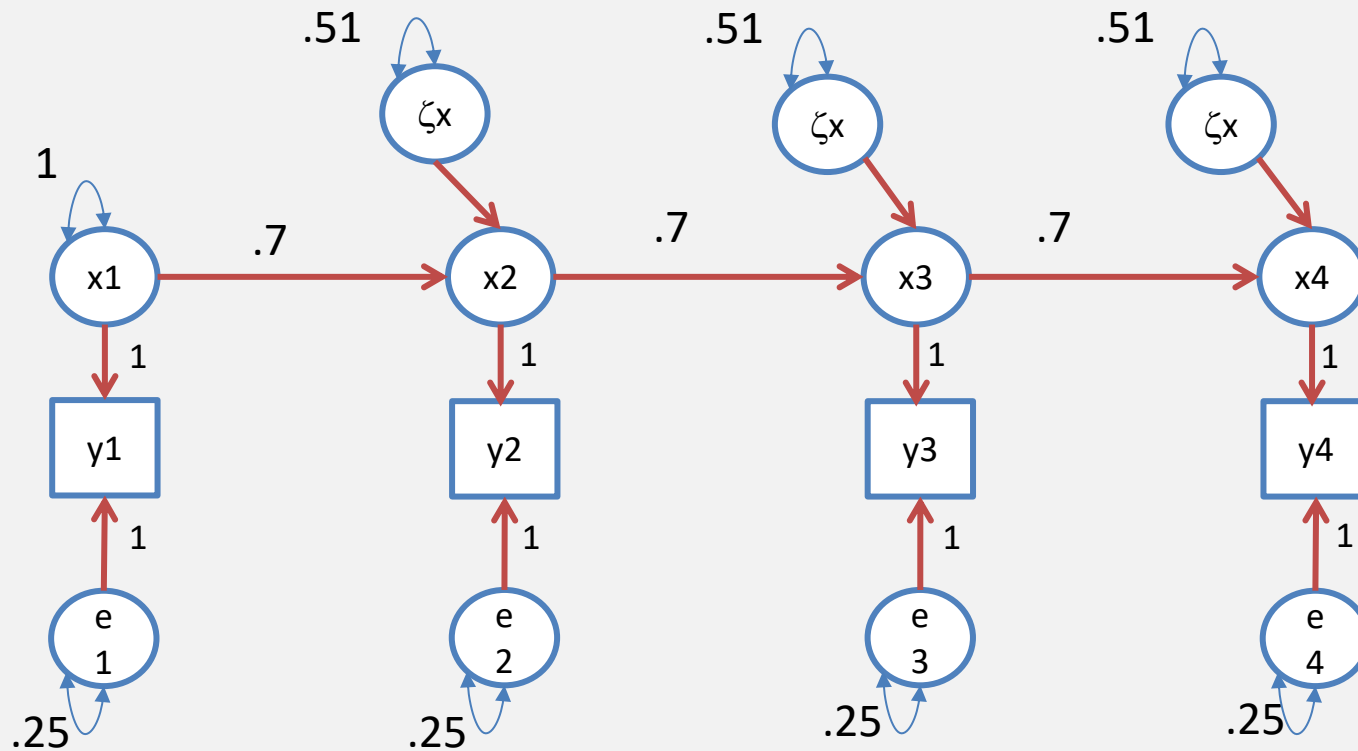
First order autoregression model.



Identification issue: $\text{var}(e_1)$ and $\text{var}(e_4)$ are not identified.

Solution set to zero, or equate :

$$\text{var}(e_1) = \text{var}(e_2) , \text{var}(e_3) = \text{var}(e_4)$$



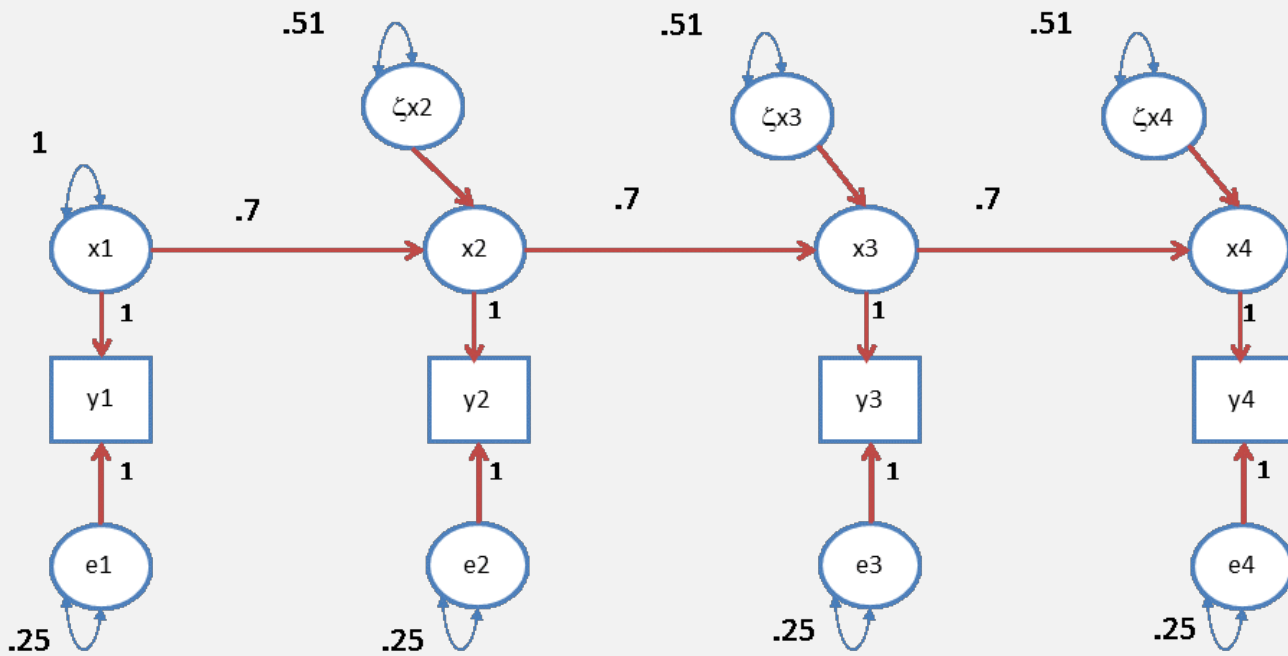
Covariance matrix

1.250	0.70	0.49	0.343
0.700	1.25	0.70	0.490
0.490	0.70	1.25	0.700
0.343	0.49	0.70	1.250

Correlation matrix

1.0000	0.560	0.392	0.2744
0.5600	1.000	0.560	0.3920
0.3920	0.560	1.000	0.5600
0.2744	0.392	0.560	1.0000

Cov y_1, y_2 : $(0.7 * 1 = 0.7$; cov $y_1, y_2 = 0.7 * 0.7 * 1 = 0.49$, etc.



reliability: $\text{rel}(x_t) = \text{var}(x_t) / \{\text{var}(x_t) + \text{var}(e_t)\}$
 $= 1 / (1 + .25) = 1 / 1.25 = .8$

$R^2: b_{t-1,t}^2 \text{var}(x_{t-1}) / \{b_{t-1,t}^2 \text{var}(x_{t-1}) + \text{var}(\zeta x_t)\}$
 $= \{.7^2 * 1\} / (.7^2 * 1 + .51) = .49 / 1 = .49$

Correlation (t,t+1) : $b_{t-1,t} \text{var}(x_{t-1}) / \{\text{sd}(y_{t-1}) * \text{sd}(y_t)\}$
 $= \{.7 * 1\} / \{\sqrt{1.25} * \sqrt{1.25}\} = .56$

These models can be applied to phenotypic, genetic and environmental timeseries.

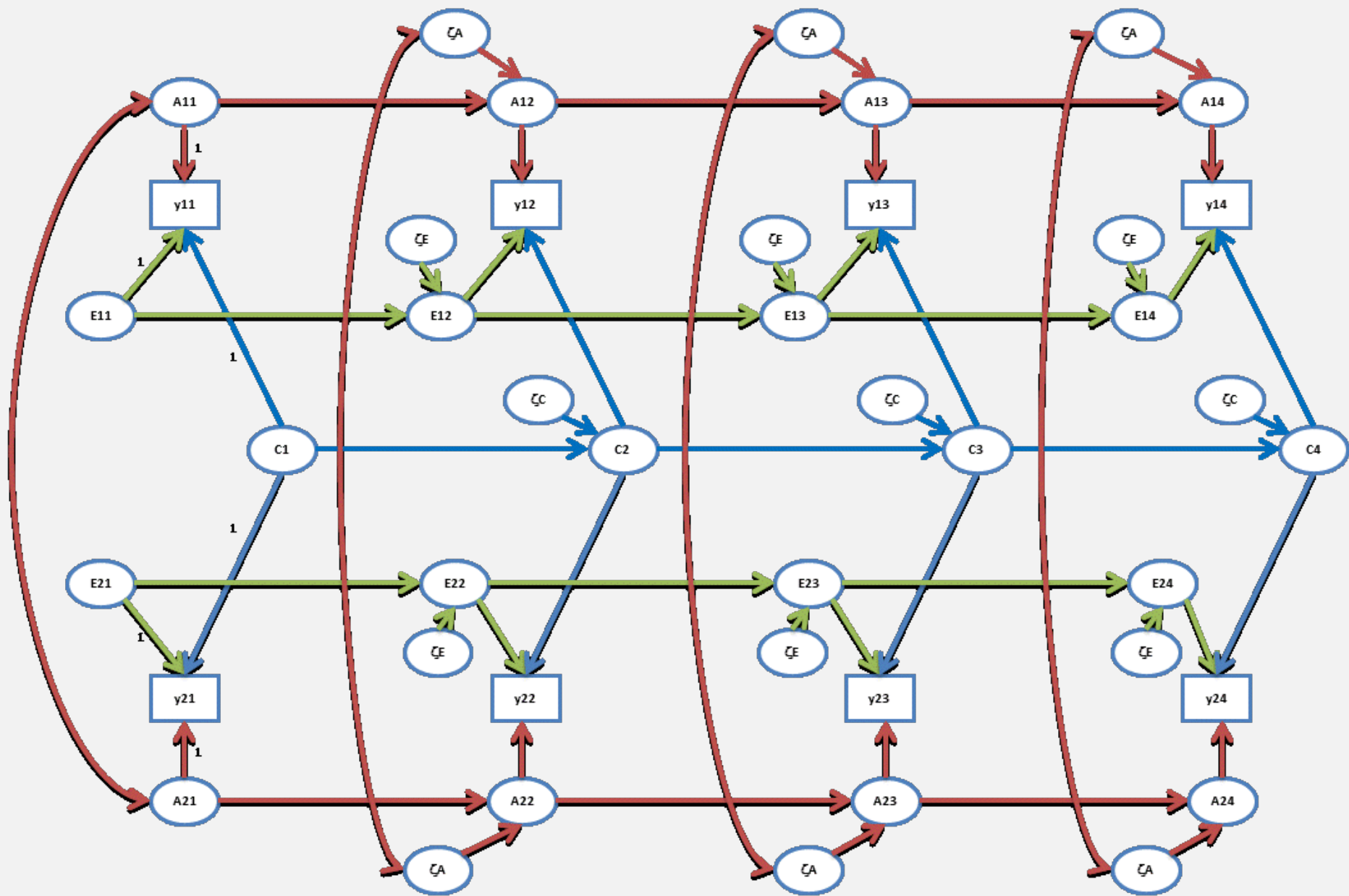
Estimation of genetic and environmental correlations

Animal / experimental studies

Bi/multivariate twin / family analyses

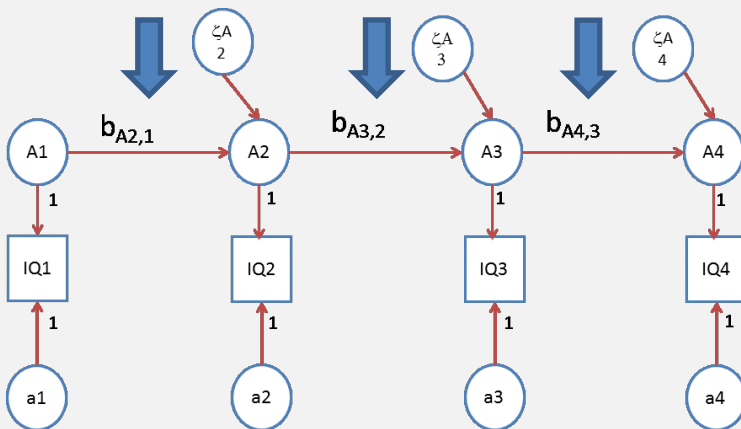
Bivariate SNP analyses

LD-score regression



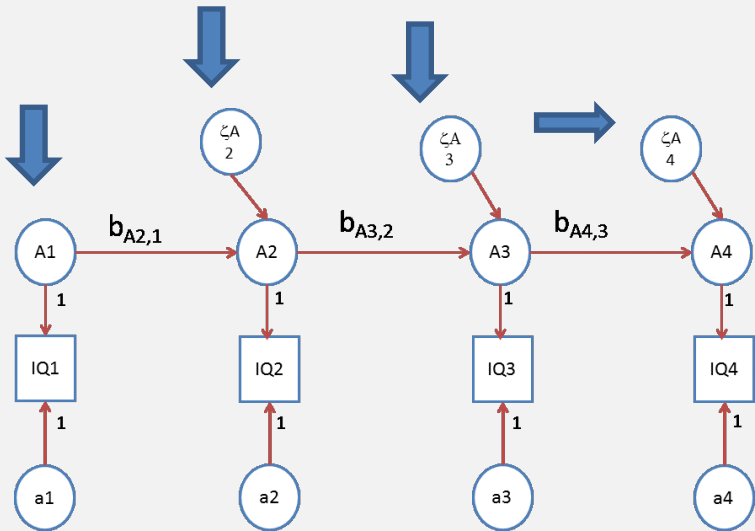
$$\Sigma_A = (I - B_A)^{-1} \Psi_A (I - B_A)^{-1} t + \Theta_A$$

$$B_A = \begin{bmatrix} 0 & 0 & 0 & 0 \\ b_{A21} & 0 & 0 & 0 \\ 0 & b_{A32} & 0 & 0 \\ 0 & 0 & b_{A43} & 0 \end{bmatrix}$$



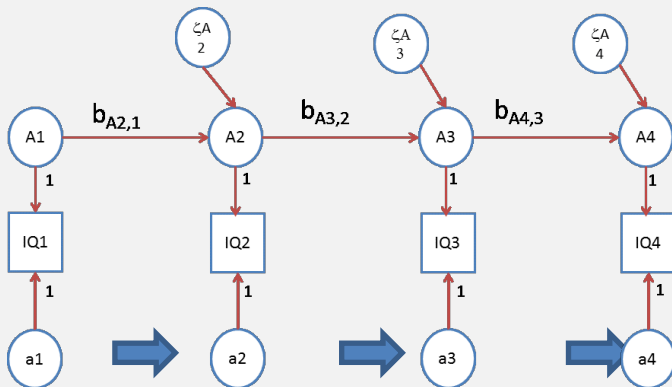
$$\Sigma_A = (I - B_A)^{-1} \Psi_A (I - B_A)^{-1 t} + \Theta_A$$

$$\Psi_A = \begin{bmatrix} \text{var}(A_1) & 0 & 0 & 0 \\ 0 & \text{var}(\zeta_{A2}) & 0 & 0 \\ 0 & 0 & \text{var}(\zeta_{A3}) & 0 \\ 0 & 0 & 0 & \text{var}(\zeta_{A4}) \end{bmatrix}$$



$$\Sigma_A = (\mathbf{I} - \mathbf{B}_A)^{-1} \Psi_A (\mathbf{I} - \mathbf{B}_A)^{-1\tau} + \Theta_A$$

$$\Theta_A = \begin{bmatrix} \text{var}(a_1) & 0 & 0 & 0 \\ 0 & \text{var}(a_2) & 0 & 0 \\ 0 & 0 & \text{var}(a_3) & 0 \\ 0 & 0 & 0 & \text{var}(a_4) \end{bmatrix}$$



Constraint on residuals, e.g.:
 $\text{var}(a_1) = \text{var}(a_2) = \text{var}(a_3) = \text{var}(a_4)$

Stability in symptoms of anxiety and depression as a function of genotype and environment: a longitudinal twin study from ages 3 to 63 years

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Genetic and Environmental Stability of Neuroticism From Adolescence to Adulthood

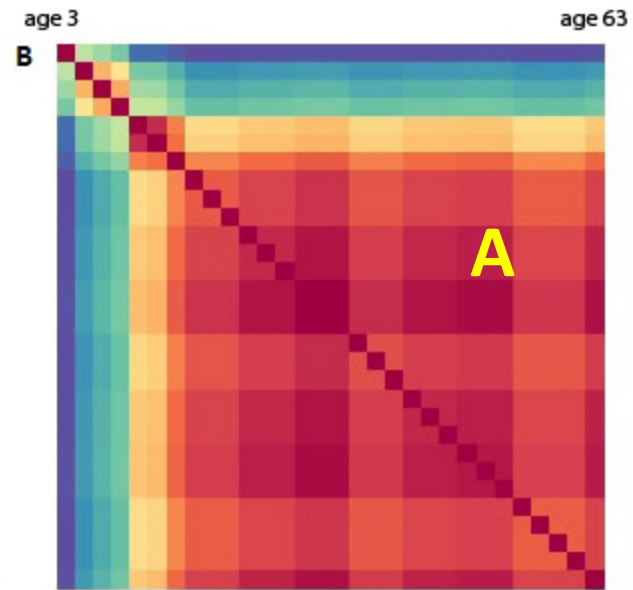
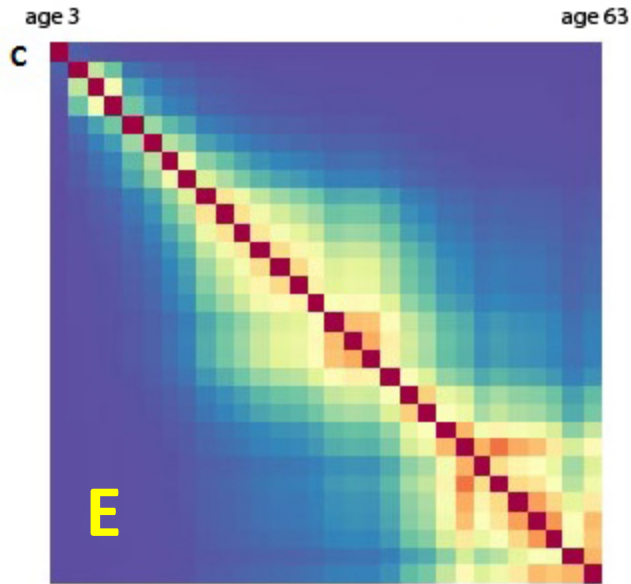
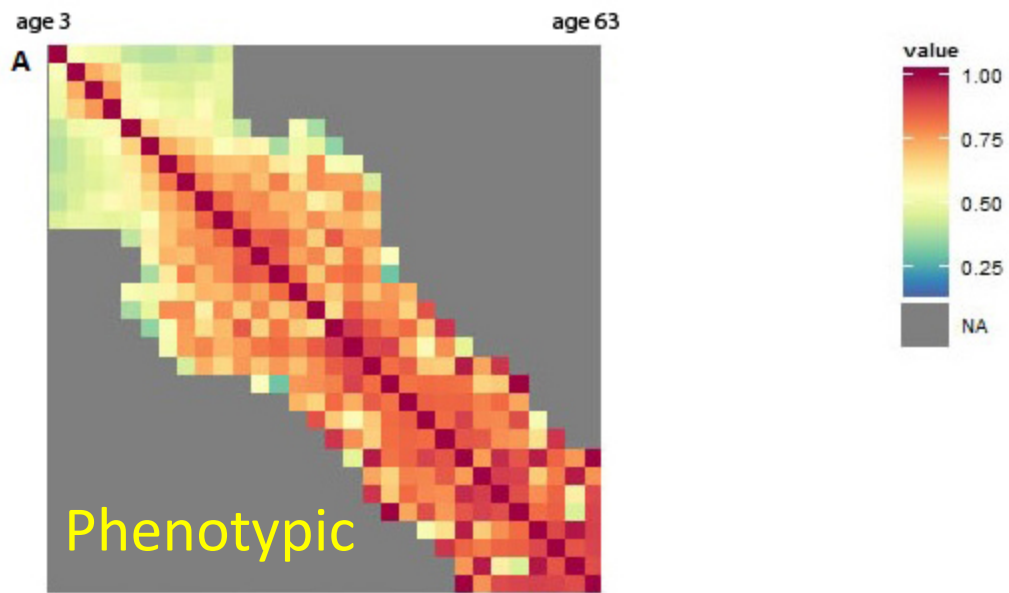
Michel G. Nivard,¹ Christel M. Middeldorp,^{1,2} Conor V. Dolan,¹ and Dorret I. Boomsma^{1,2,3}
¹Department of Biological Psychology, VU University Amsterdam, the Netherlands
²Neuroscience Campus Amsterdam, Amsterdam, the Netherlands
³EMGO⁺ Institute for Health and Care Research, Amsterdam, the Netherlands

Genetic and Environmental Stability in Attention Problems Across the Lifespan: Evidence From the Netherlands Twin Register

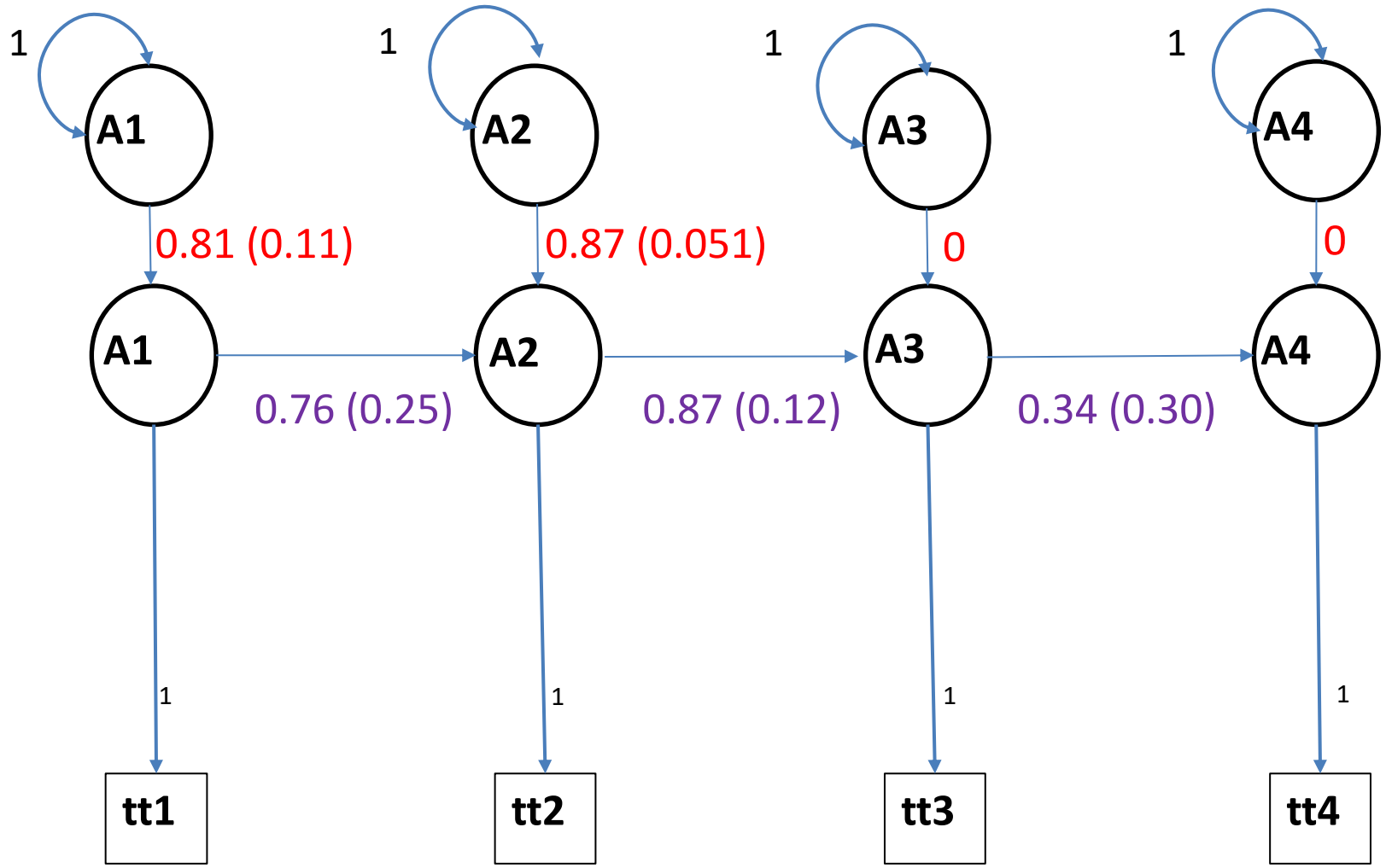
Kees-Jan Kan, Ph.D., Conor V. Dolan, Ph.D., Michel G. Nivard, M.Sc., Christel M. Middeldorp, Ph.D., Catharina E.M. van Beijsterveldt, Ph.D., Gonneke Willemsen, Ph.D., Dorret I. Boomsma, Ph.D.

This stability model can be applied at the genetic and non-genetic level.

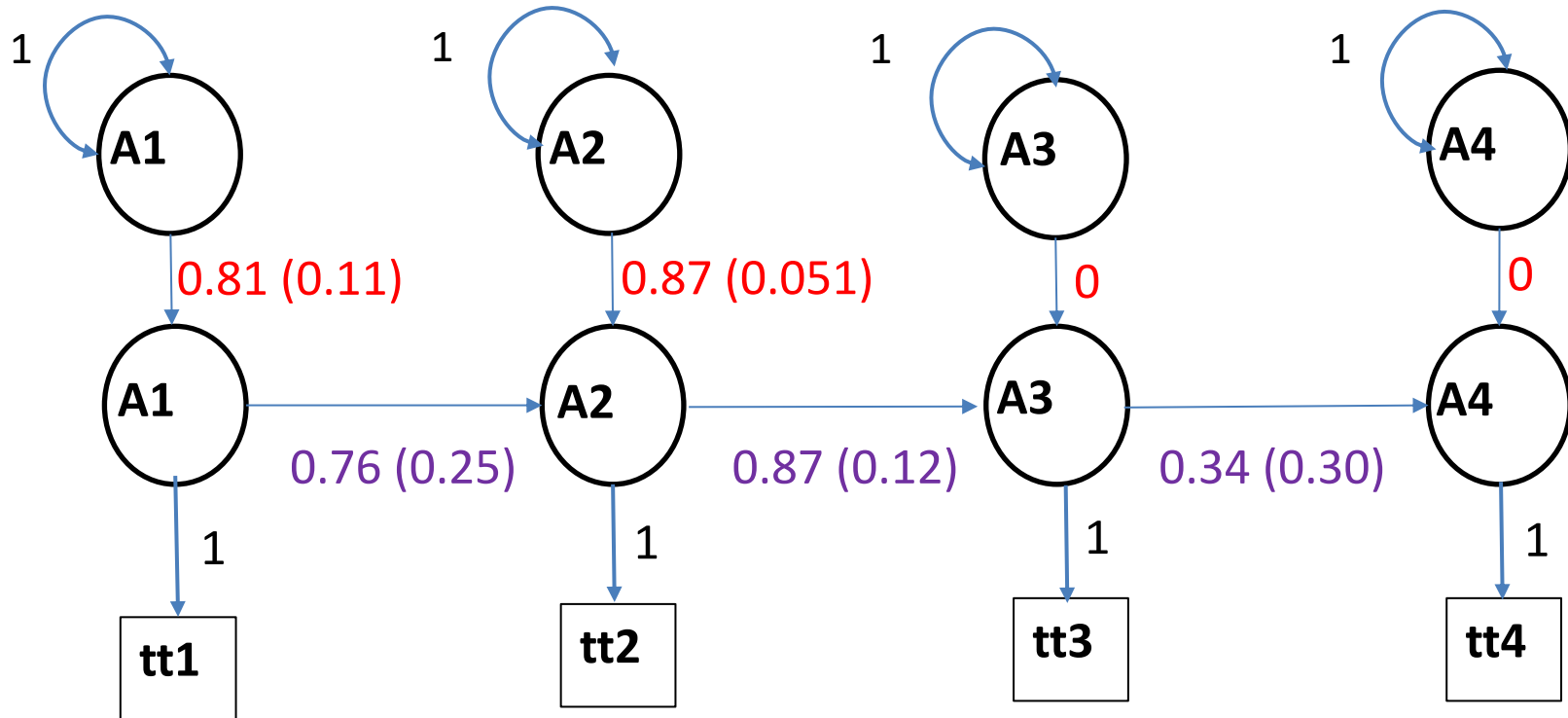
e.g. Nivard et al, 2014



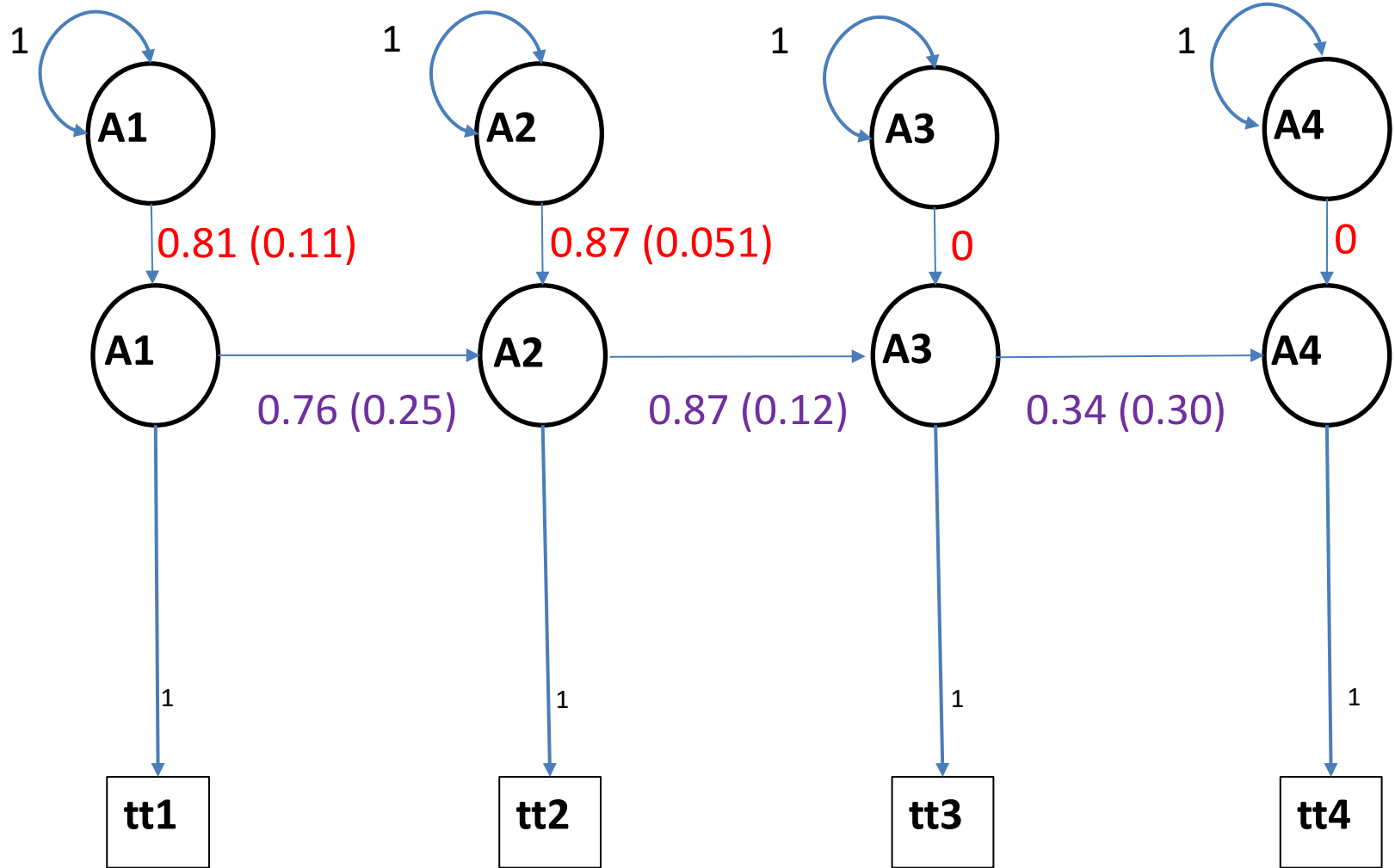
Anx/depression stability due to A and E from 3y to 63 years



Genetic Simplex (type D personality). Latent factors in circles, twin-time data in squares. Loadings from latent to observed traits and variation of innovations constrained at 1. Correlations between twins are specified at the level of the innovations (1 for MZ and 0.5 for DZ pairs).



Innovation genetic variance				
Variance due to Genetic Transmission				
Total $V(A)$				



A Innovation	0.81²=0.66	0.87²=0.76	0	0
Transmission	-	0.66*0.76=0.50	1.26*0.87=1.10	1.10*0.34
Total V(A)	0.66	1.26	1.10	0.37

We will look at several approaches in an analysis of BMI data in young twins from the Netherlands Twin Register.

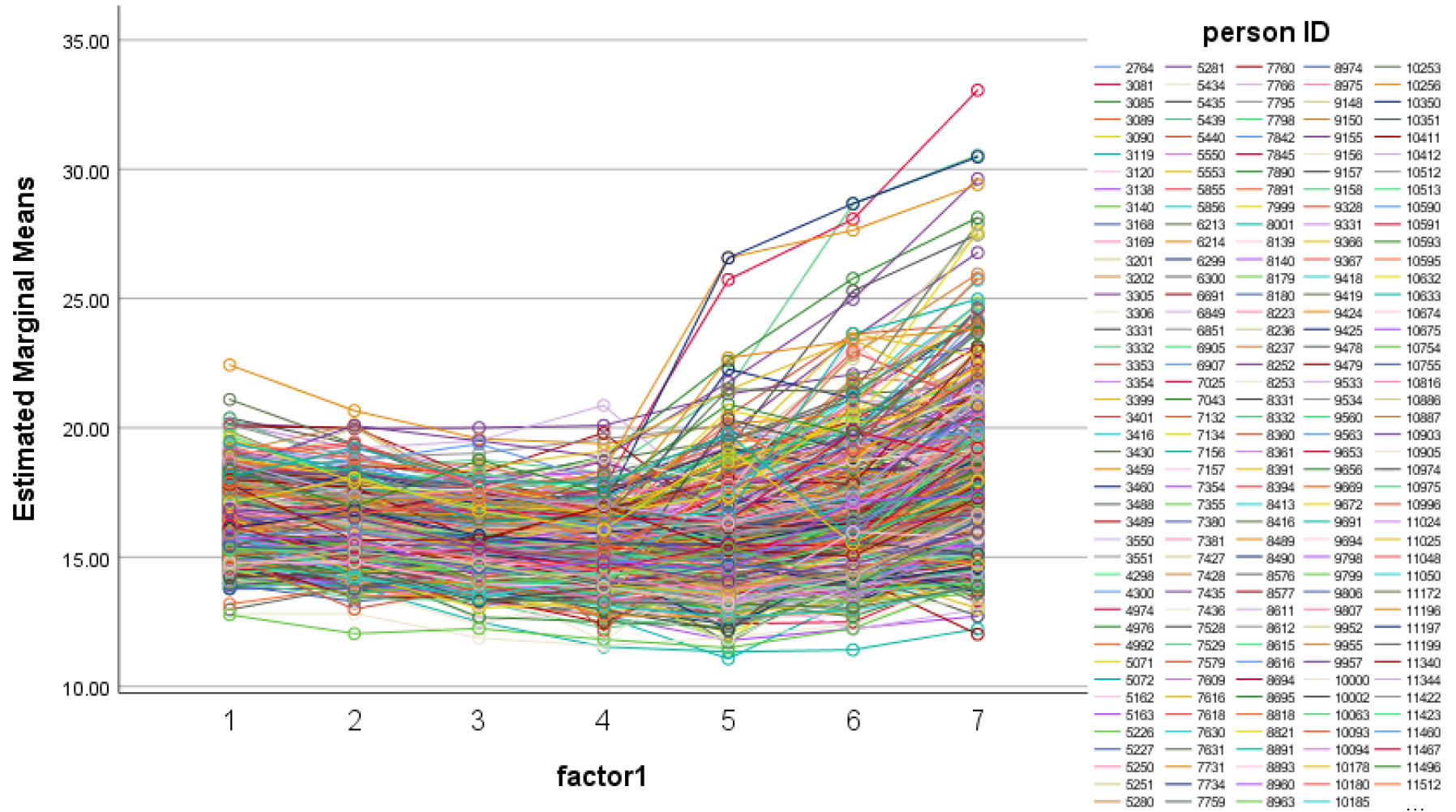
- 1) Saturated model
- 2) Cholesky 'model' (decomposition)
- 3) Simplex model

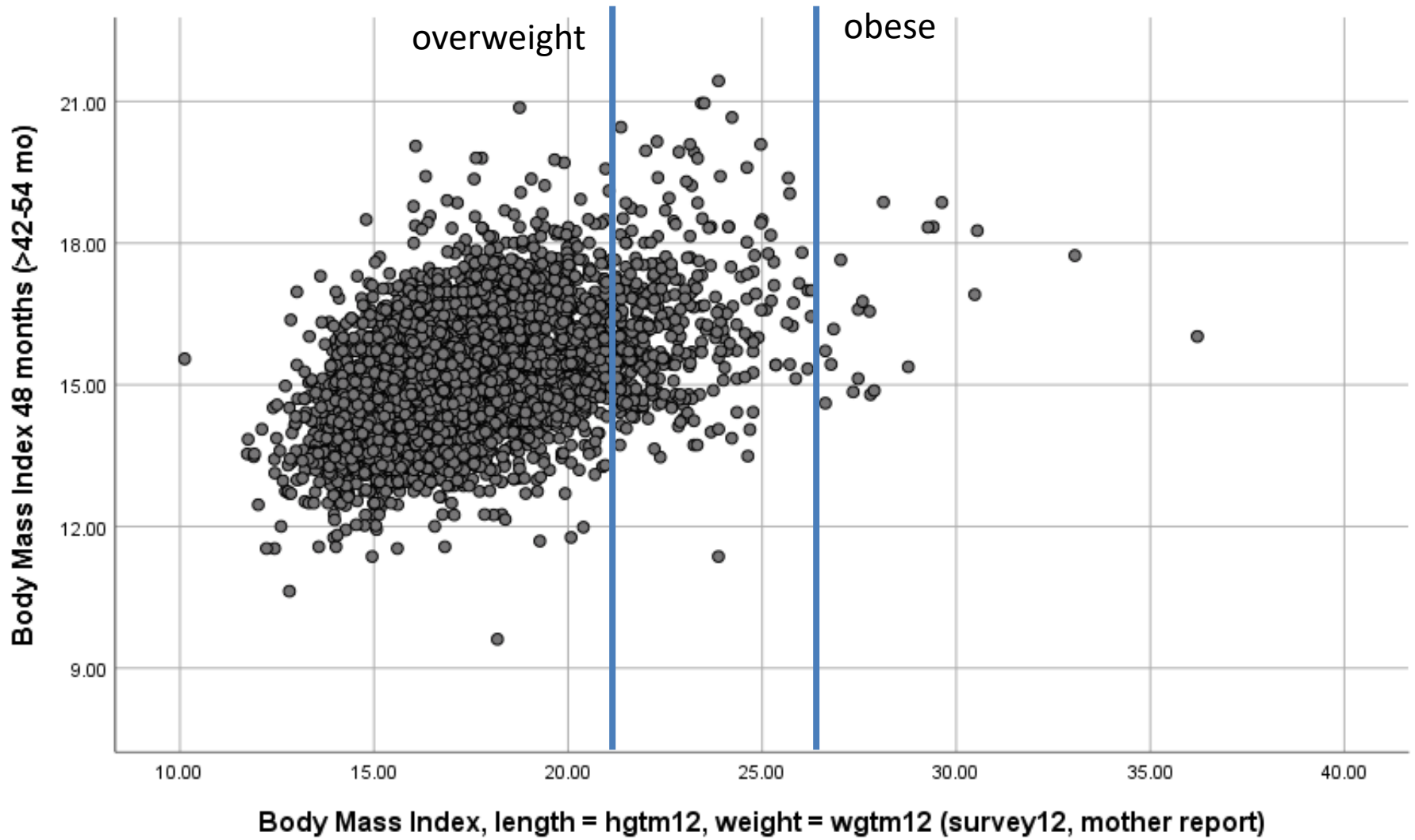
The BMI data set has data on 35,120 kids starting at birth.

We selected kids with complete data only and will analyze BMI at age 4, 7, 9 and 12 years.

For 7 ages: 1.5, 2, 3, 4, 7, 10 and 12 years (N= 734): individual data

Estimated Marginal Means of MEASURE_1



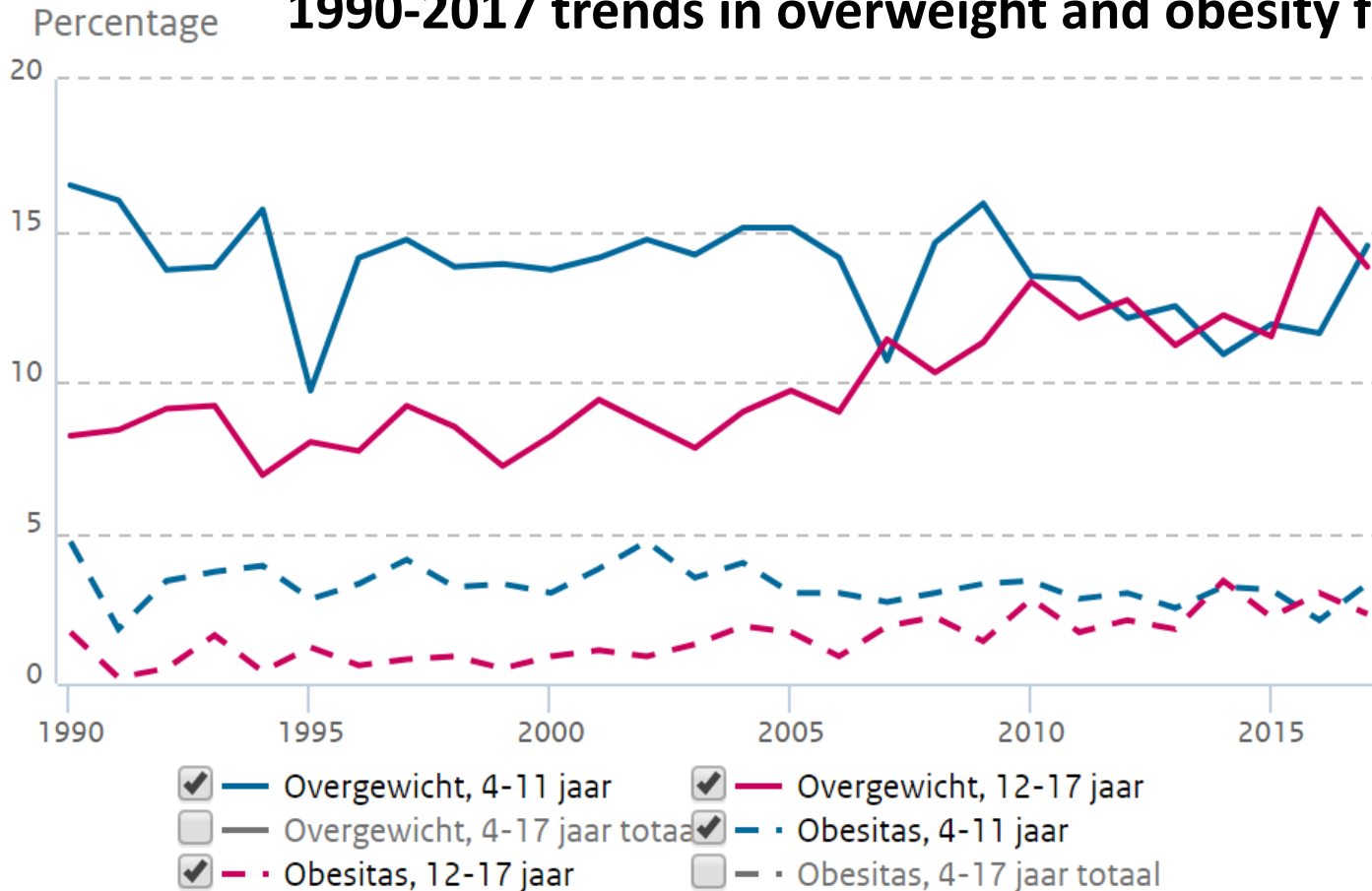


Correlations

BMI across age (bold: all data / below diagonal : complete data)

BMI 4 yr	1	.580	.493	.431
BMI 7 yr	.590	1	.729	.658
BMI 10 yr	.515	.743	1	.785
BMI 12 yr	.442	.670	.803	1

1990-2017 trends in overweight and obesity for kids 4-18 year



Bron: CBS Gezondheidsenquête (tot en met 2013); daarna

Gezondheidsenquête/Leefstijlmonitor CBS i.s.m. RIVM

- De gegevens zijn [gestandaardiseerd](#) naar de Nederlandse bevolking van 2017
- De [BMI-grenswaarden van overgewicht en obesitas van kinderen en jongeren](#) wijken af van die van volwassenen, zie [Definities](#).

QUIZ

1) you have > 1 observation per person (from twin pairs) and fit a saturated model: what is the output / information ?

2) on the same data you fit a Cholesky decomposition: what is the information contained in the output?

3) if you aim for a model which recognizes time dependency in data what are the options?



André-Louis Cholesky

QUIZ

1) > 1 observation per person (from twin pairs) and fit a saturated model: what is the output / information ?

a) ??????????????????

b) ...

c) ...

QUIZ

1) > 1 observation per person (from twin pairs) and fit a saturated model: what is the output / information ?

- a) ??????????????????
- b) An estimate for the mean value of the traits in twins
- c) An estimate of the correlation for all traits in (MZ and DZ) pairs
- d) An estimate of the phenotypic correlation between traits
- e) An estimate of the (MZ and DZ) cross-correlations between traits
- f) All of the above (except a))

QUIZ

2) on the same data you fit a Cholesky 'model' (decomposition): what is the information contained in the output?

- a) The phenotypic correlations
- b) The genetic correlations
- c) The environmental correlations
- d) b and c