
Extending Simplex model to model $Ph \rightarrow E$ transmission

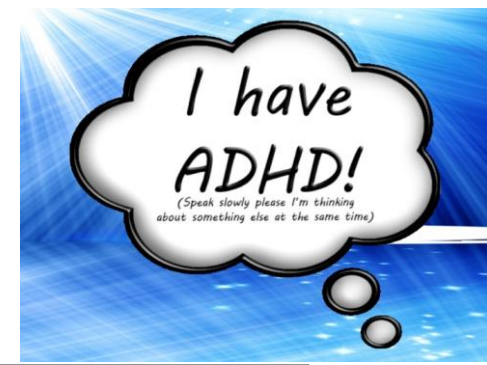
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Nature versus Nurture

Role GEcov in Psychopathology



ADHD diagnosis

GE covariance: Genotypic control over environmental exposures

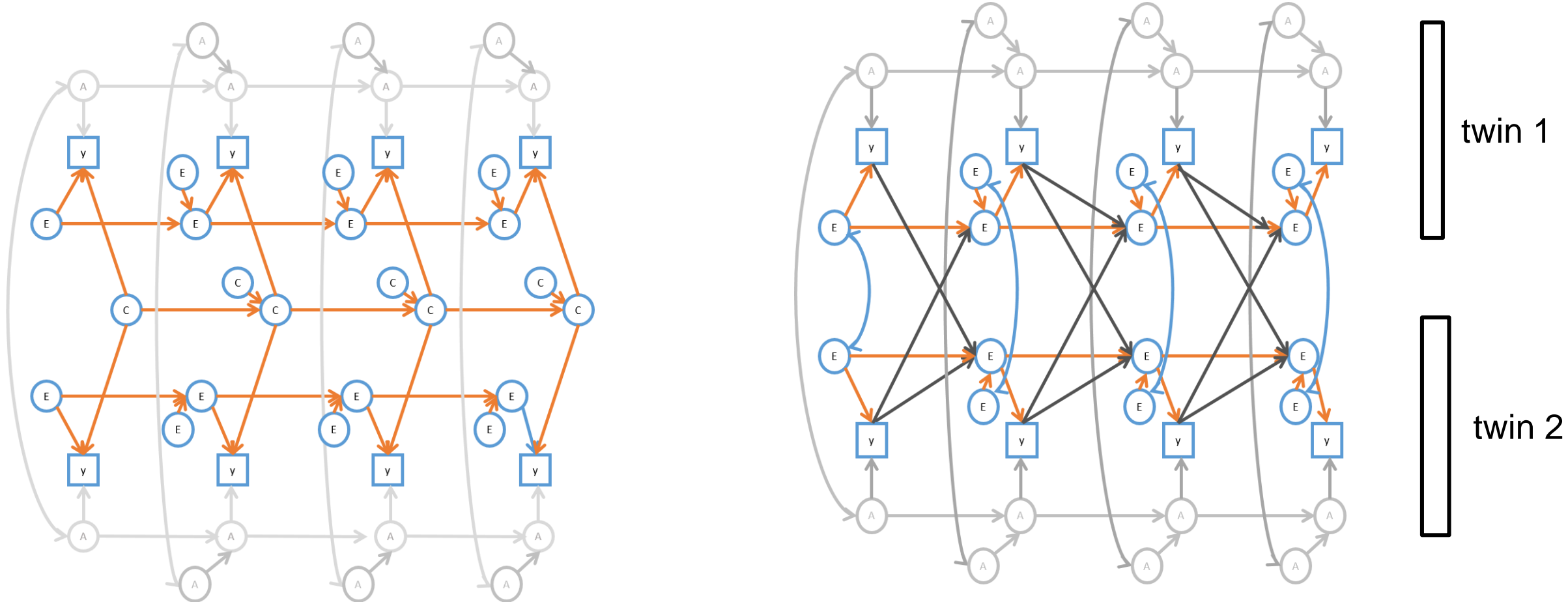
- Evocative:** Behavior of individual evokes reaction from environment consistent with genotype
- Do people with ADHD evoke different behavior in others?**
 - **ADHD child :** Intrudes on games more often
 - **ADHD adolescent :** Has more frequent school disciplinary actions
 - **ADHD adult:** more often has work conflicts
- Active:** Individuals actively seek out environments consistent with their phenotype (e.g. Niche Picking)
- Do people diagnosed with ADHD seek more risky environments**
 - **ADHD child :** climb more trees
 - **ADHD adolescent :** does more binge drinking and has casual sex more often
 - **ADHD adults:** has more driving violations, uses more drugs

Translate path diagram into matrices

ACE SIMPLEX

Ph->E SIMPLEX

Competing models



OBJECTIVE: model the observed covariance matrix

S = 4 observed variables for twin 1, 4 for twin 2, thus 8 x 8

twin 1				twin 2			
t = 1	t = 2	t = 3	t = 4	t = 1	t = 2	t = 3	t = 4
varP11							
covP11P12	varP12						
covP11P13	covP12P13	varP13					
covP11P14	covP12P14	covP13P14	varP14				
covP11P21	covP12P21	covP13P21	covP14P21	varP21			
covP11P22	covP12P22	covP13P22	covP14P22	covP21P22	varP22		
covP11P23	covP12P23	covP13P23	covP14P23	covP21P23	covP22P23		
varP23							
covP11P24	covP12P24	covP13P24	covP14P24	covP21P24	covP22P23		
covP23P24	varP24						



Matrices used in Model

$$\Sigma = \Lambda(I-B)^{-1}\Psi(I-B)^{-t}\Lambda^t + \Theta$$

ne = number of latent variables in the model

ny = number of observed variables

Σ = Sigma = Expected covariance matrix observed variables y
= ny x ny

Λ = Lambda = Factor Loading Matrix = ny x ne

B = Beta = Matrix with regression coefficients between **latent variables**
= ne x ne

Θ = Theta = Matrix with residuals = ny x ny

Ψ = Psy = Matrix with variances and covariance between **latent variables**
= ne x ne

Dimensions of matrices for 4 time points in **normal** simplex specification

$$\Sigma MZ = \Lambda * \text{solve}(I - B) * \Psi MZ * t(\text{solve}(I - B)) * t(\Lambda) + \Theta MZ$$

$$\Sigma DZ = \Lambda * \text{solve}(I - B) * \Psi DZ * t(\text{solve}(I - B)) * t(\Lambda) + \Theta DZ$$

▪ **Twins = 2**

▪ **Time points = 4**

▪ **ny = 4 time points for 2 twins = 8**

▪ **ne = latent variables A, C, E at each timepoint for each twin: 3 x 4 = 12**

Σ = Sigma	=	8 x 8	= ny x ny
Λ = Lambda	=	8 x 24	= ny x ne
B = Beta	=	24 x 24	= ne x ne
Ψ = Psy	=	24 x 24	= ne x ne
Θ = Theta	=	8 x 8	= ny x ny

Does not allow us to regress E on P

PROBLEM:

Regression from Observed to Latent variable

Normally

PSY matrix only has latent variables.

In case of twin modeling: the variables A, C, and E → 24 x 24

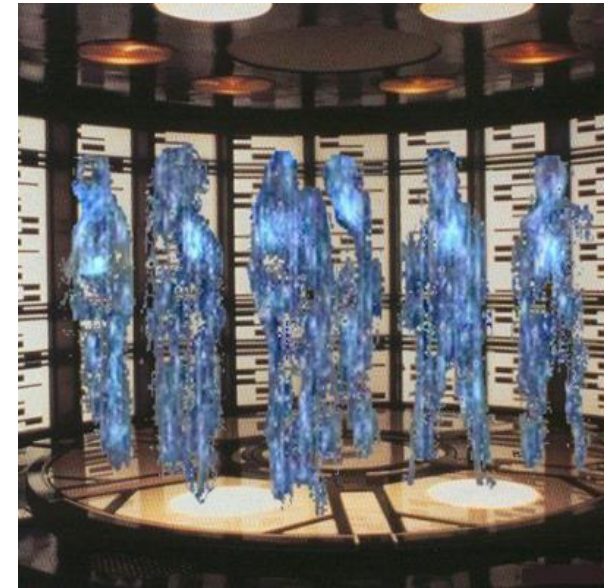
Problem

Impossible to regress from observed to latent

For that we need all variables in Ψ , thus P, A, C, and E.

Solution: Change dimensions of matrices

Beam observed variables into Ψ space



Dimensions of matrices for 4 time points in **our** simplex specification

$$\Sigma MZ = \Lambda * \text{solve}(I - B) * \Psi MZ * t(\text{solve}(I - B)) * t(\Lambda) + \Theta MZ$$

$$\Sigma DZ = \Lambda * \text{solve}(I - B) * \Psi DZ * t(\text{solve}(I - B)) * t(\Lambda) + \Theta DZ$$

▪ Twins = 2

▪ Time points = 4

▪ ny = 4 time points for 2 twins = 8

▪ ne = latent variables A, C, E plus observed P at each time point for each twin: **4 x**

4 x 2 = 32

$\Sigma = \text{Sigma} = 8 \times 8 = ny \times ny$

$\Lambda = \text{Lambda} = 8 \times 32 = ny \times ne$

$B = \text{Beta} = 32 \times 32 = ne \times ne$

$\Psi = \text{Psy} = 32 \times 32 = ne \times ne$

$\Theta = \text{Theta} = 8 \times 8 = ny \times ny$

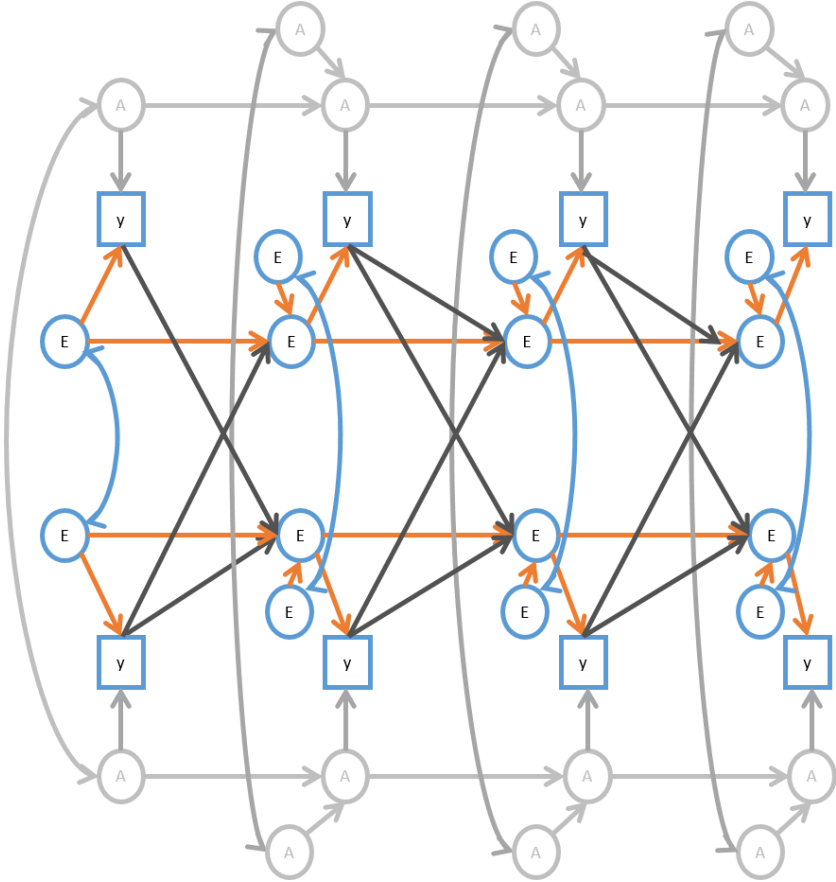
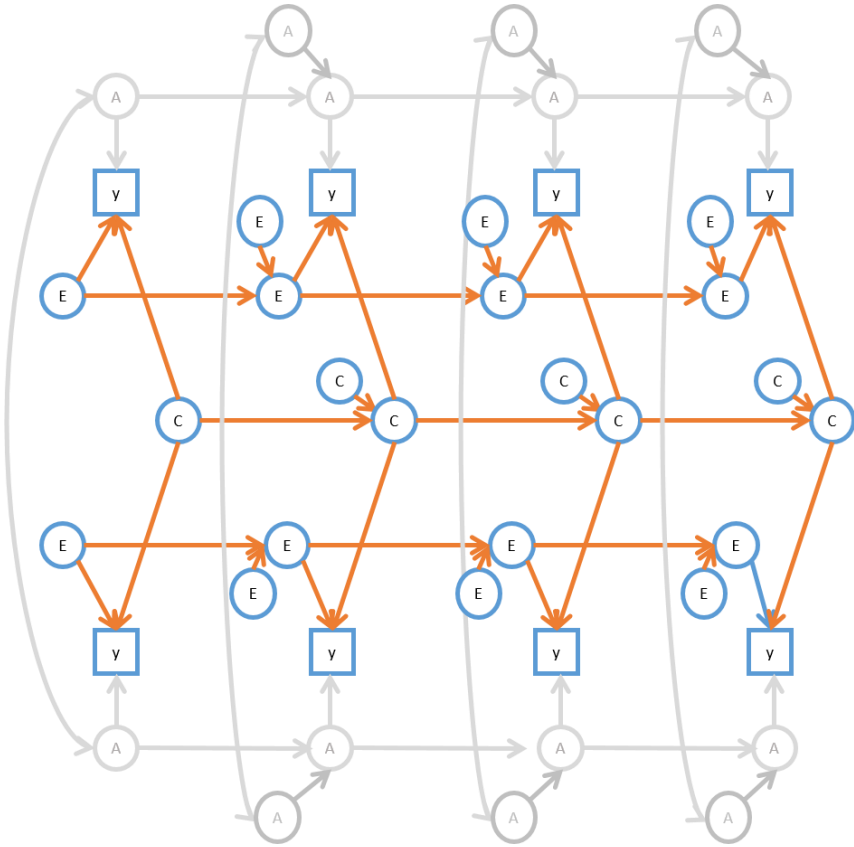
SIMPLEX
matrix explanation

Specification SIMPLEX in OpenMx



General code

Open:
SIMPLEX practical



PRACTICAL

Replicate findings in following article

J. *Intell.* 2014, 2, 1-x manuscripts; doi:10.3390/jintelligence20x000x

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Article

Can GE-covariance originating in phenotype to environment transmission account for the Flynn effect?

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Abstract: The Dickens and Flynn model of the Flynn effect (generational increases in mean IQ) assigns an important role to genotype-environment covariance (GE-cov). We quantify GE-cov in a longitudinal simplex model by modeling it as phenotype to environment (Ph->E) transmission in twin data. The model fits as well as the standard genetic simplex model, which assumes uncorrelated genetic and environmental influences. We use the results to explore numerically the possible role of GE-cov in amplifying increases in environmental means. Given the estimated Ph->E transmission parameters, GE-cov resulted in an amplification of 1.57 (full scale IQ) to 1.7 (perforal IQ). The results lend credence to the role of GE-cov in the Flynn effect.

Keywords: genotype-environment covariance; Flynn effect; Dickens and Flynn model; longitudinal genetic modeling.

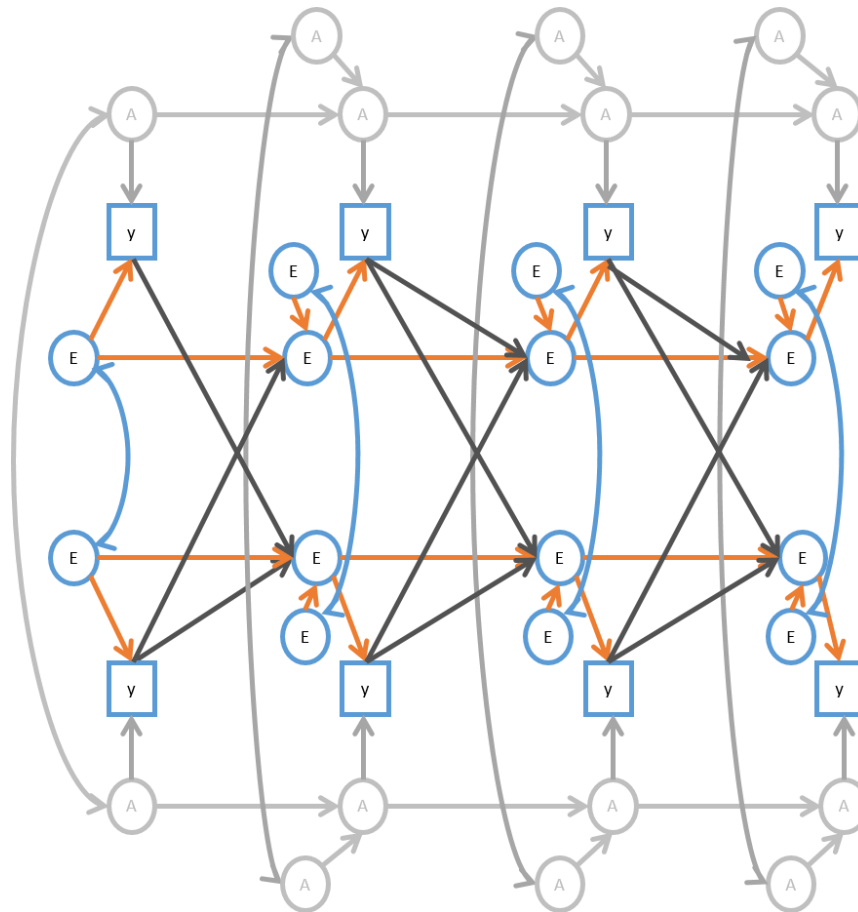
1. Introduction

8266 WORDS ENGLISH (UNITED STATES)

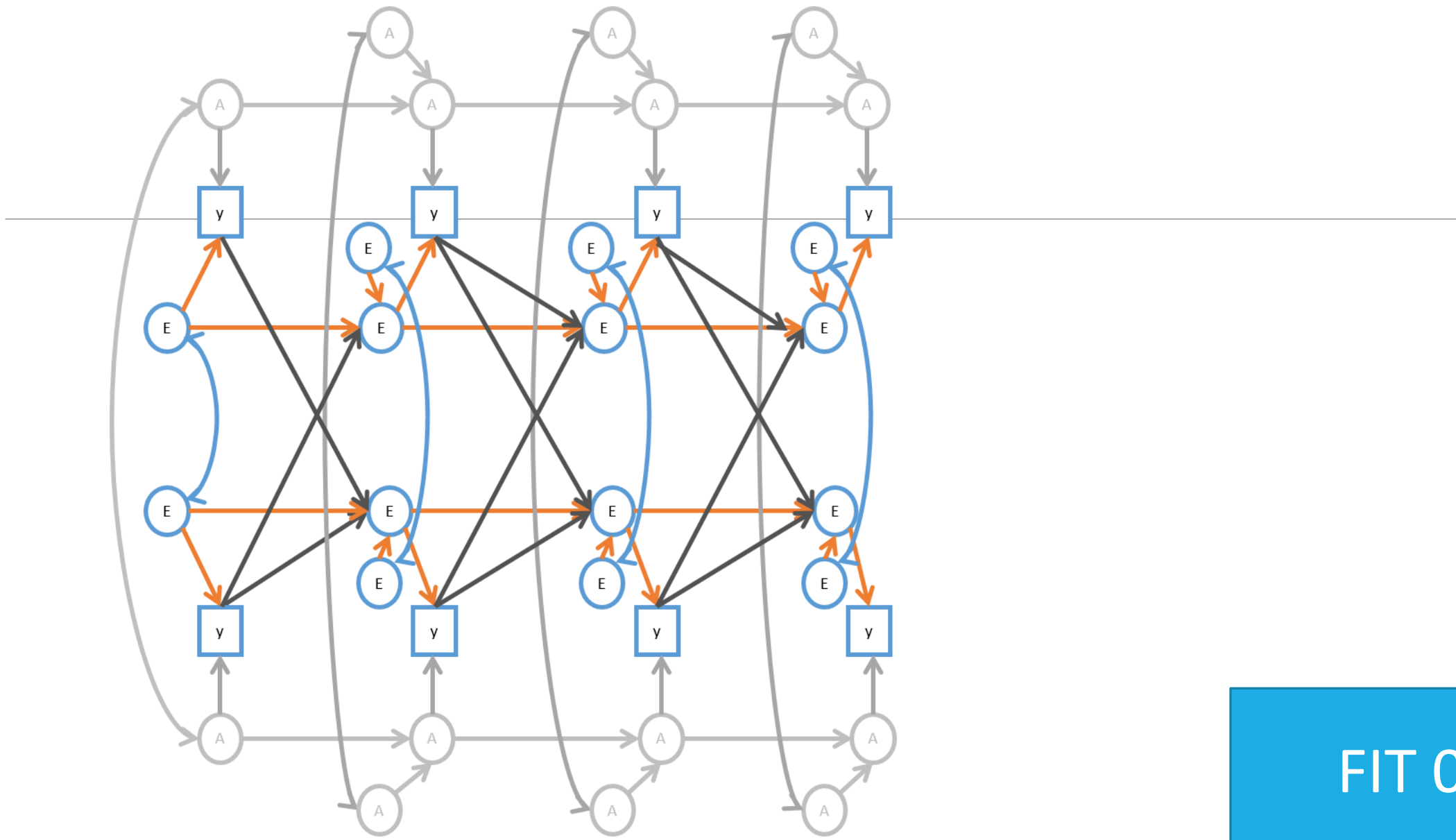
EN 8:56 6-3-2014

Model fitting: Ph \rightarrow E SIMPLEX MODEL

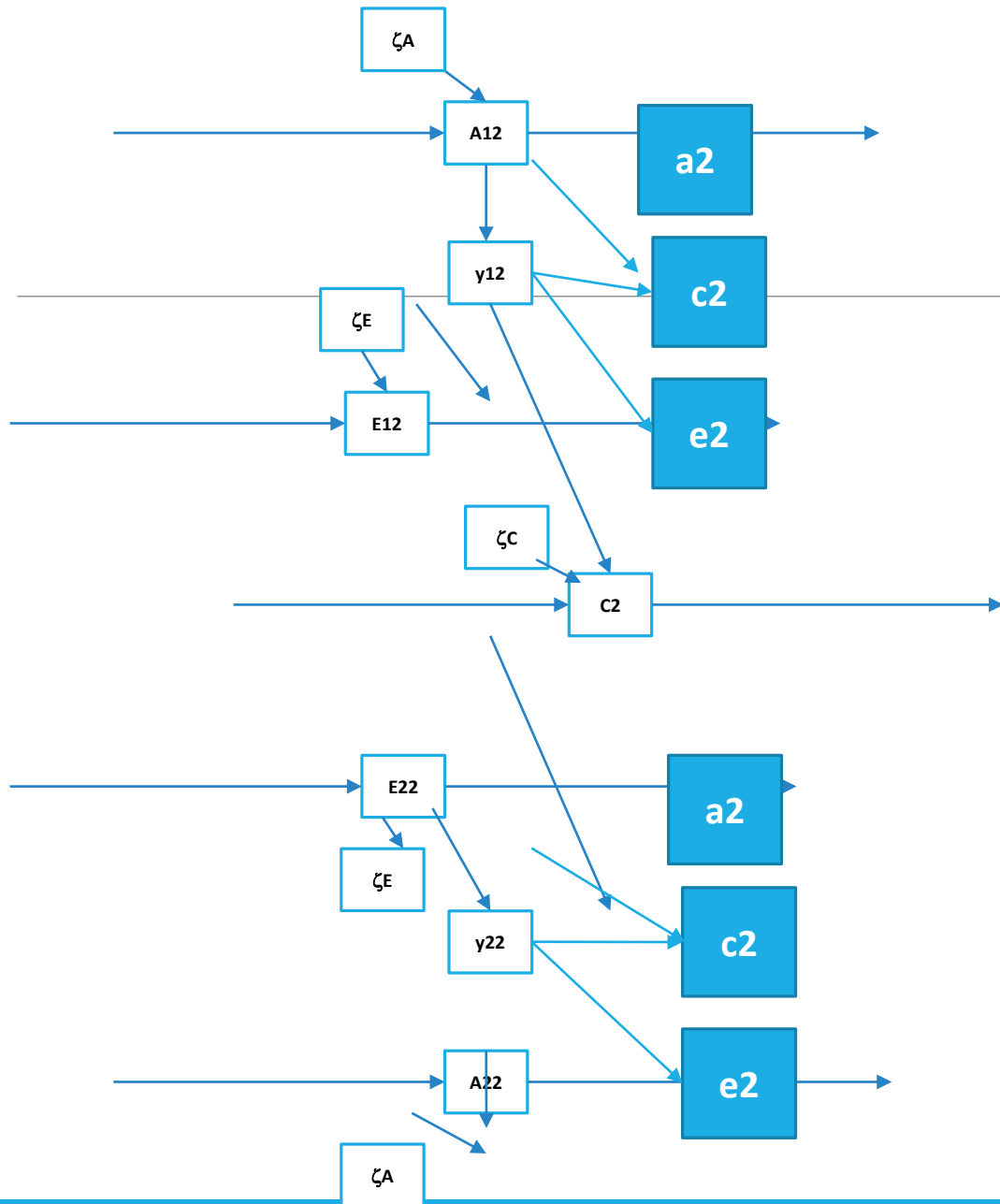
DATA: Full scale Intelligence
In MZ and DZ twins
Measured 4 time points
NTR data



Return to:
SIMPLEX practical

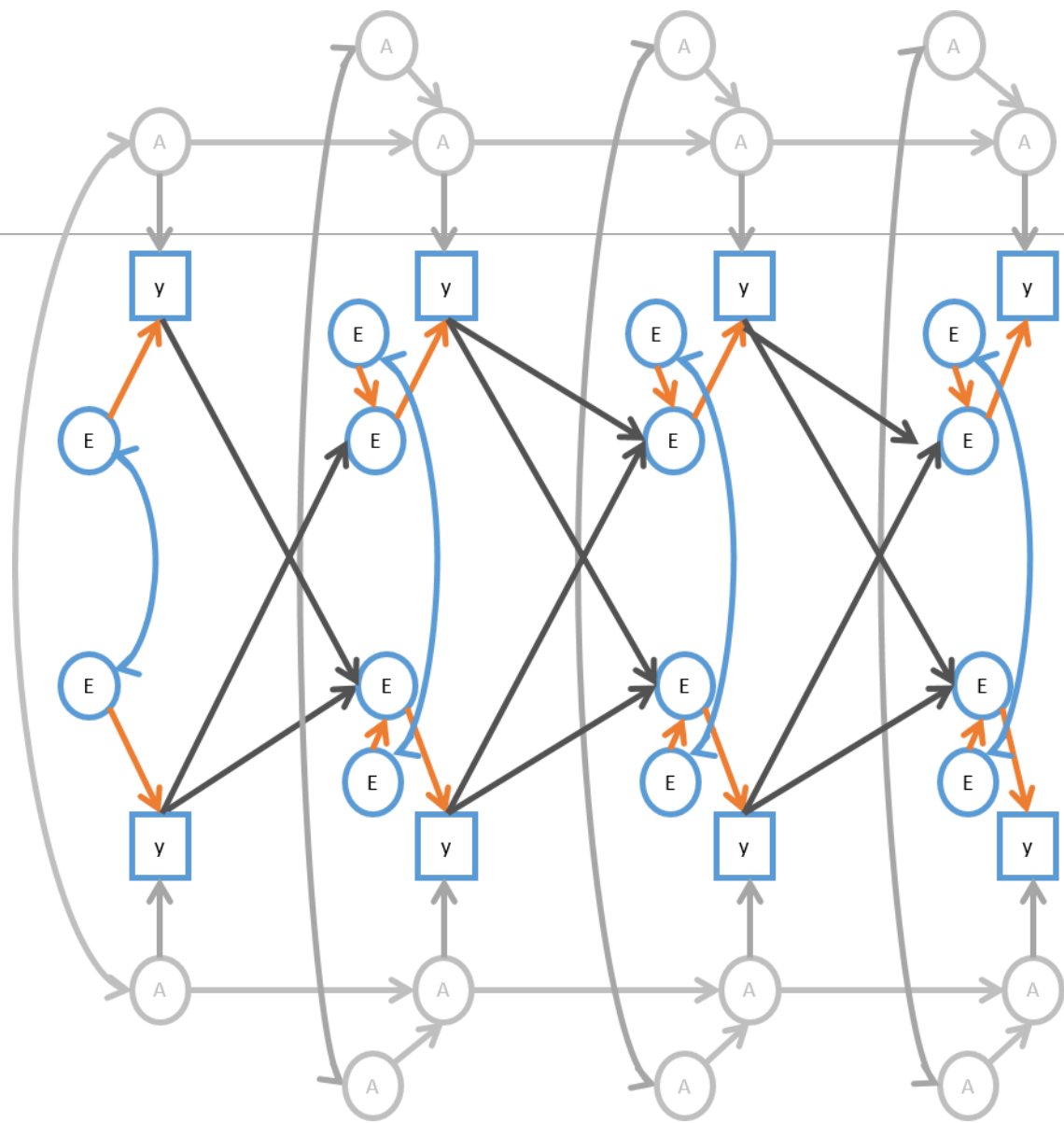


FIT 0



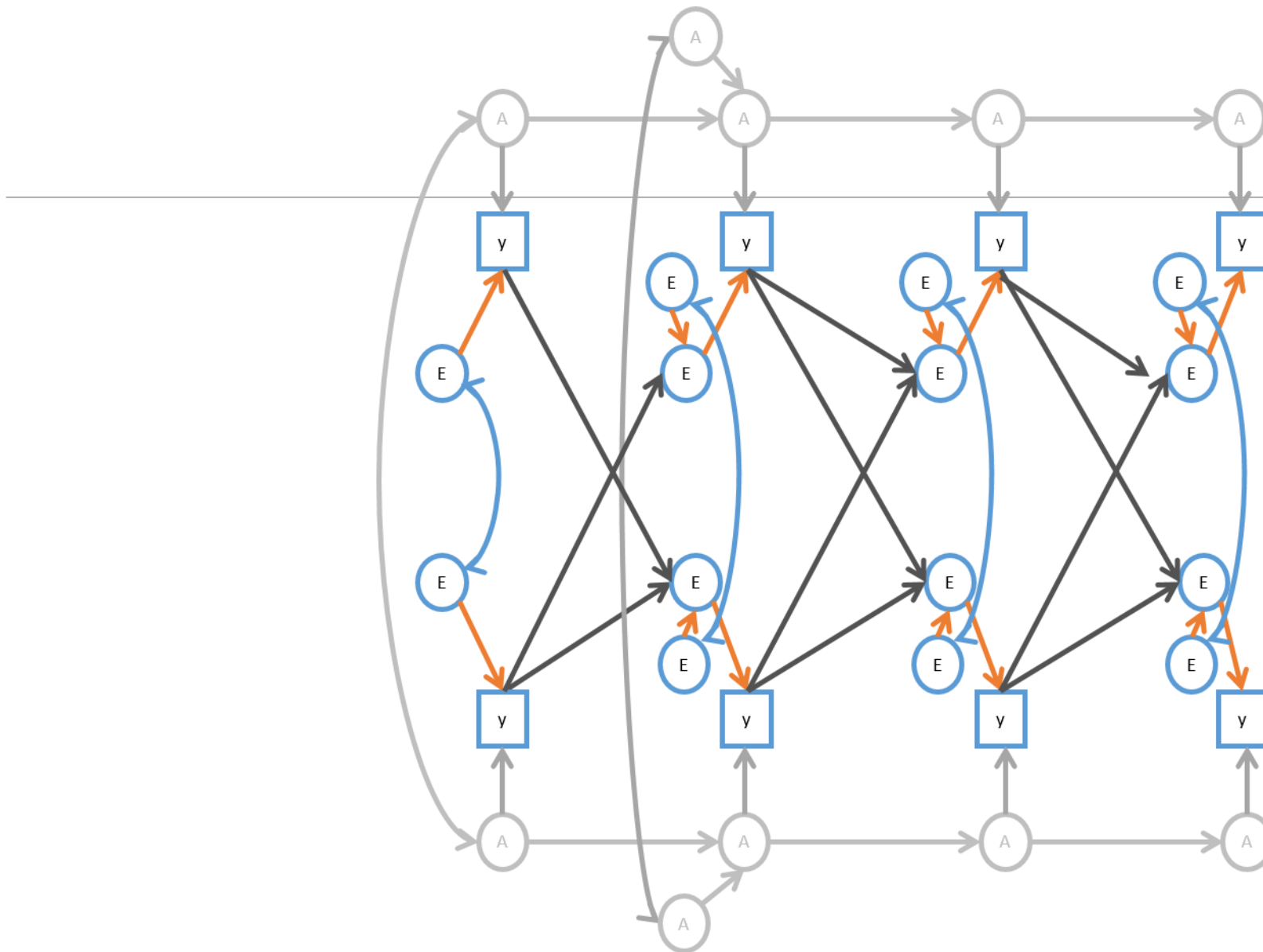
DROP a time specific

FIT 1



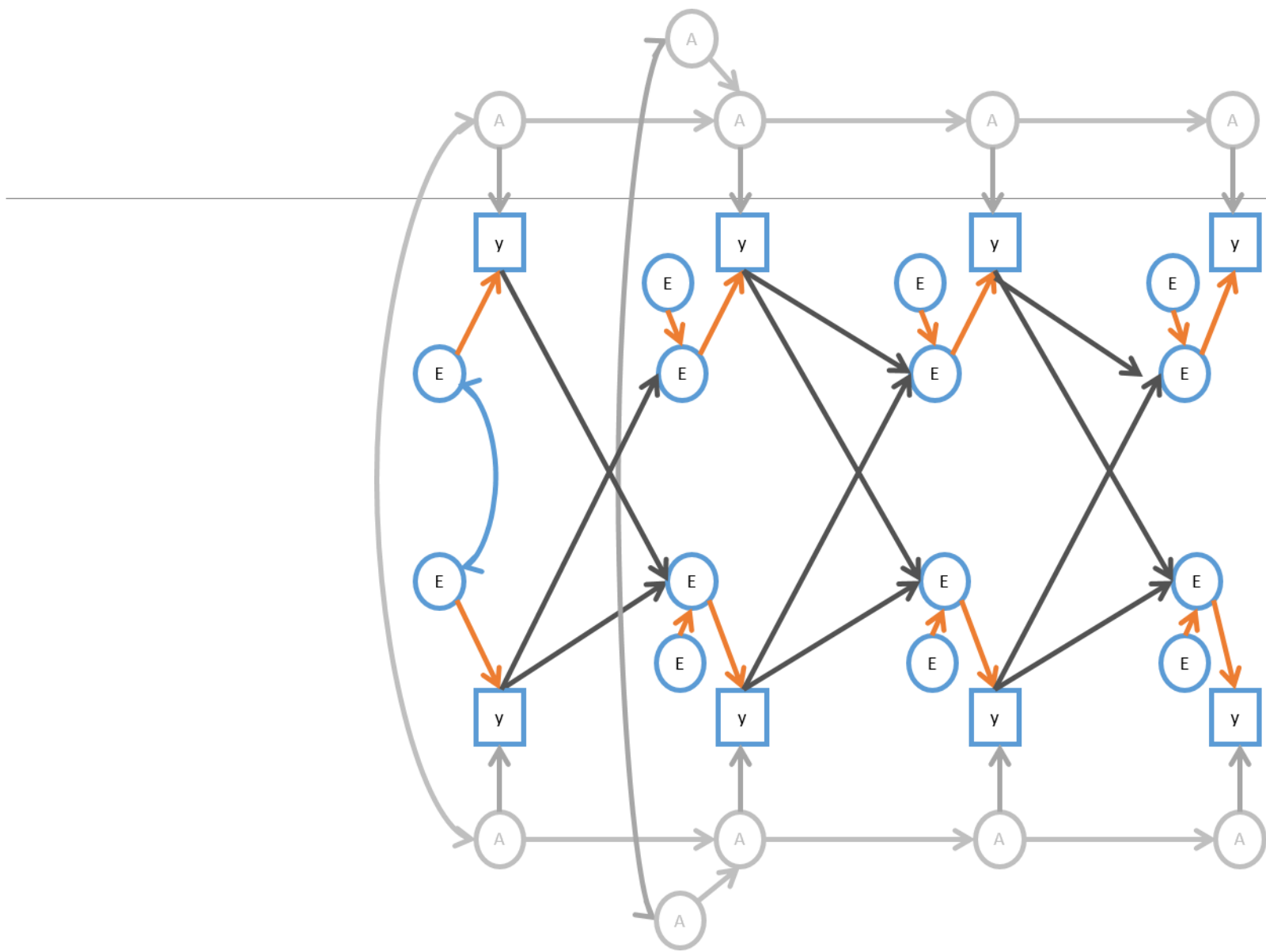
DROP E
TRANSMISSION

FIT 2



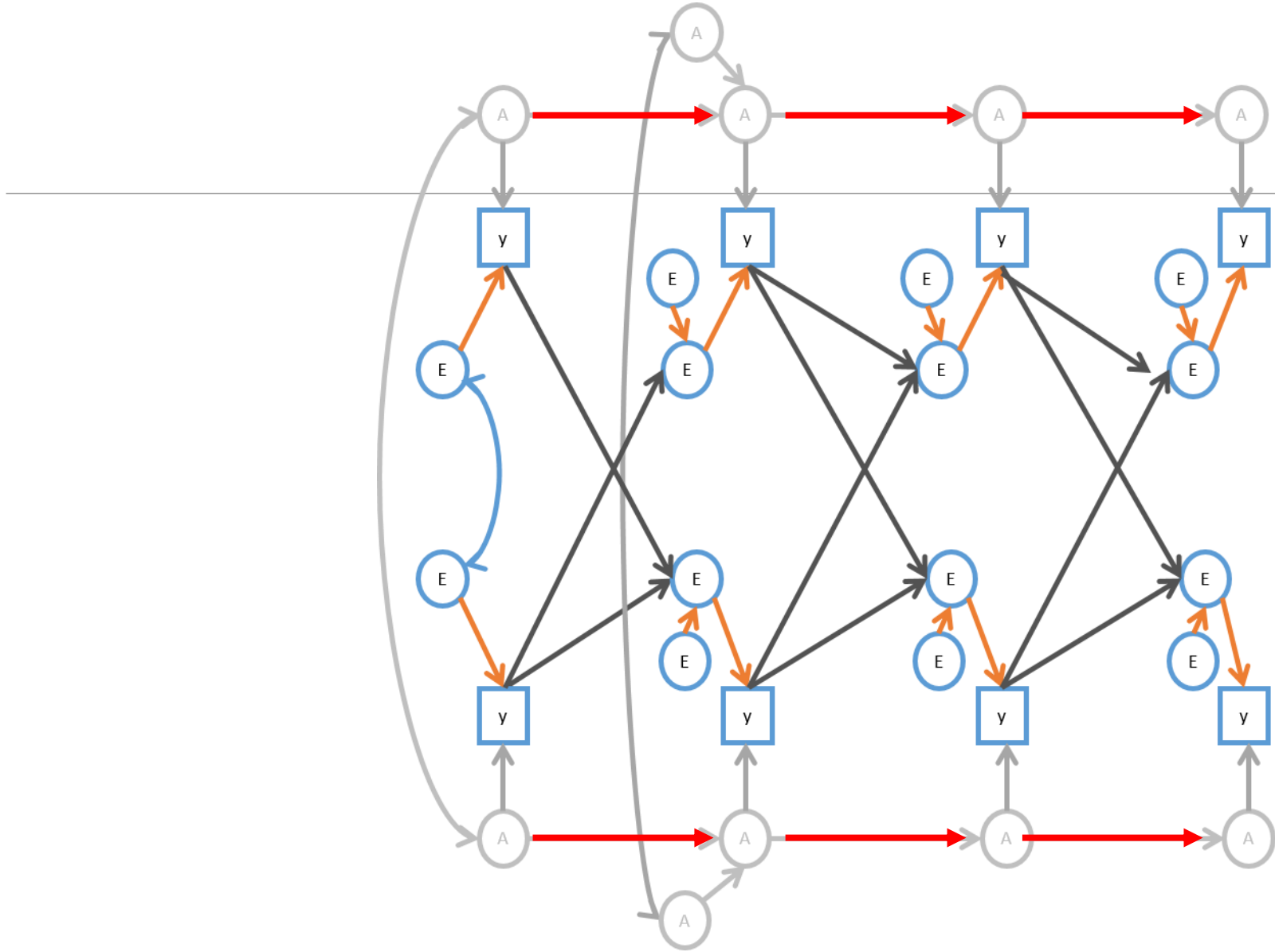
DROP A
 innovation
 Time 3 and time 4

FIT 3



DROP
covariance
between E

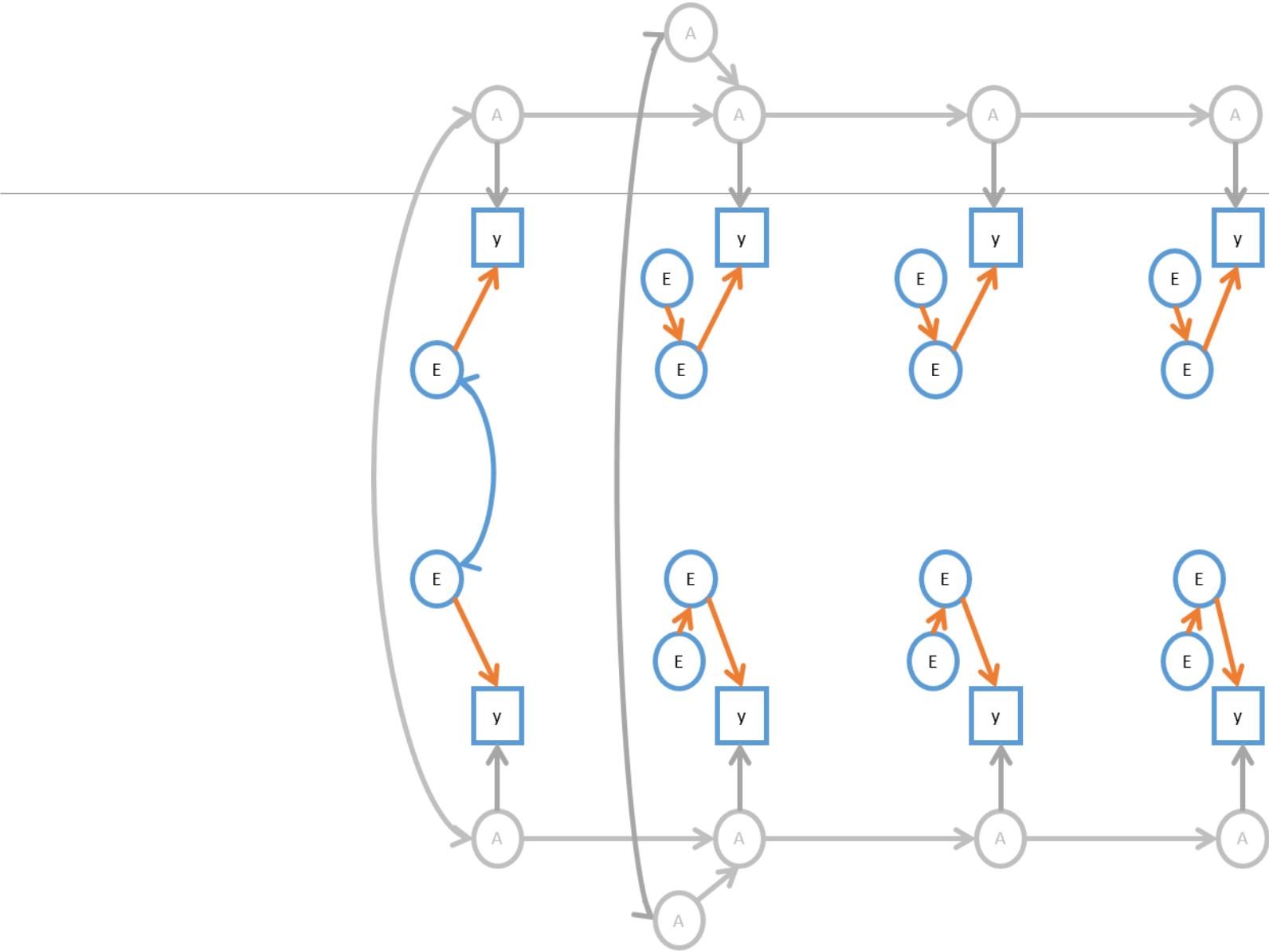
FIT 4



SET bA equal
over time

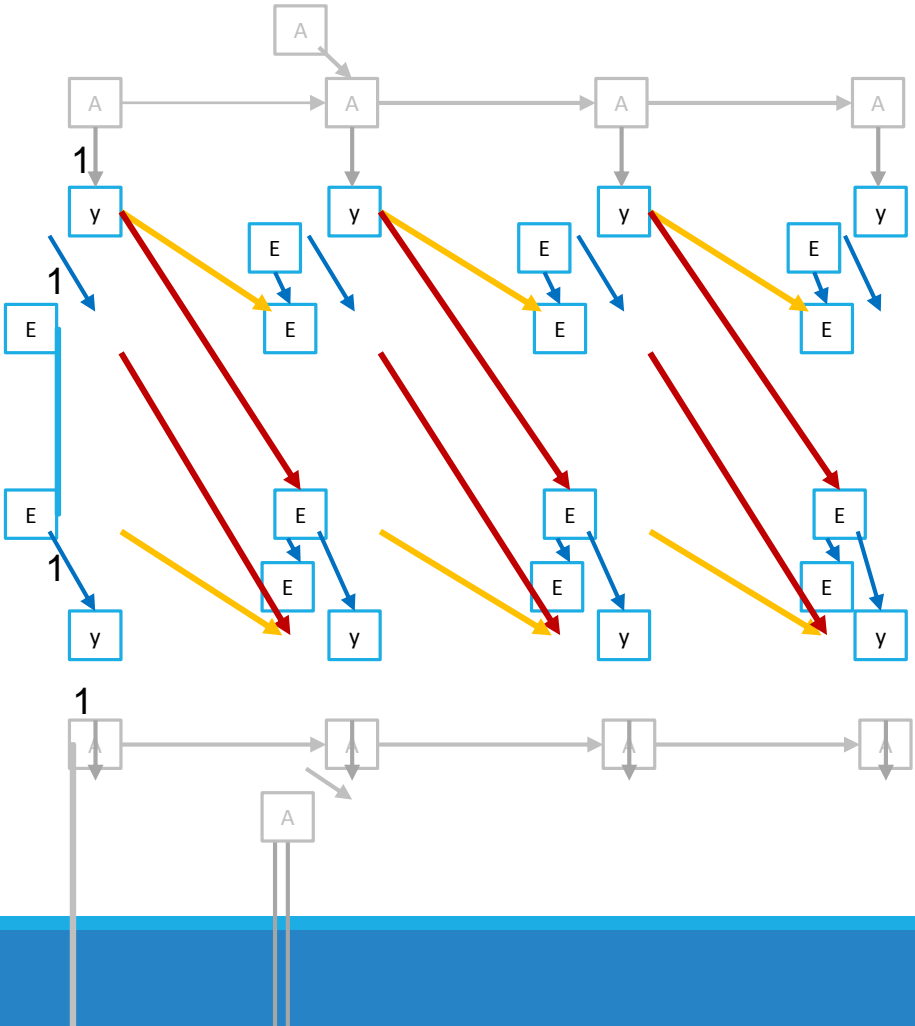
FIT 5

TEST PH- \rightarrow E
transmission



FIT 6

Final model



A1

E1

within

1			
0.86	1		
0.86	1.00	1	
0.86	1.00	1.00	1
0.00	0.00	0.00	0.00
0.35	0.30	0.30	0.30
0.50	0.53	0.53	0.53

G-E covariance

1			
0.41	1		
0.20	0.43	1	
0.09	0.30	0.47	1

A1

E1

Correlated E – But not C!

0.53	0.59	0.59	0.59
0.50	0.43	0.43	0.43
0.43	0.50	0.50	0.50
0.43	0.50	0.50	0.50
0.43	0.50	0.50	0.50

A2

between

0.00	0.00	0.00	0.00
0.22	0.19	0.19	0.19
0.33	0.35	0.35	0.35
0.36	0.40	0.40	0.40

E2

0.63	0.3	0.1	0.0
0.36	0.2	0.2	0.2
0.18	0.2	0.3	0.3
0.09	0.2	0.3	0.3