

UNIVERSITY OF AMSTERDAM Faculty of Social and Behavioural Sciences

One-stage meta-analytic structural equation modeling

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Content

- 1. What is (MA)SEM?
- 2. Univariate versus multivariate methods for MASEM
- 3. Simulation study results
- 4. Discussion

Confirmatory technique to fit hypothesized models to data

Evaluate indirect effects, relations between latent and observed variables, use fit-indices to compare competing models

E.g. Factor models, path models, full SEM

No raw data needed (covariance matrix and sample size are sufficient)

Example of a path model: Theory of planned behavior from Ajzen and Fishbein (1980)



Example of a factor model



The covariance matrix between the observed variables is modeled as a function of SEM parameters

	V1	V2	V3	V4
V1	σ_{11}			
V2	σ_{21}	σ_{22}		
V3	σ_{31}	σ_{32}	σ ₃₃	
V4	σ_{41}	σ_{42}	σ_{43}	σ_{44}



Ψ_{11} Ψ_{21}	Ψ22		
$\beta_{31}\psi_{11} + \beta_{32}\psi_{21}$	$\beta_{31}\psi_{21} + \beta_{32}\psi_{22}$	$ \begin{pmatrix} \beta_{31} \psi_{11} + \beta_{32} \psi_{21} \end{pmatrix} \beta_{31} + \\ \begin{pmatrix} \beta_{31} \psi_{21} + \beta_{32} \psi_{22} \end{pmatrix} \beta_{32} + \psi_{33} \\ \end{pmatrix} $	
$\begin{array}{l} \beta_{31}\beta_{43}\psi_{11} + \\ \beta_{32}\beta_{43}\psi_{21} \end{array}$	$\begin{array}{l} \beta_{31}\beta_{43}\psi_{21}\text{+}\\ \beta_{32}\beta_{43}\psi_{22} \end{array}$	$\begin{aligned} & \left(\beta_{31}\beta_{43}\psi_{11}+\beta_{32}\beta_{43}\psi_{21}\right)\beta_{31}+\\ & \left(\beta_{31}\beta_{43}\psi_{21}+\beta_{32}\beta_{43}\psi_{22}\right)\beta_{32}+\\ & \beta_{43}\psi_{33} \end{aligned}$	$\begin{aligned} & (\beta_{31}\beta_{43}\psi_{11}+\beta_{32}\beta_{43}\psi_{21})\beta_{31}\beta_{43}+\\ & (\beta_{31}\beta_{43}\psi_{21}+\beta_{32}\beta_{43}\psi_{22})\beta_{32}\beta_{43}+\\ & \beta_{43}{}^2\psi_{33}+\psi_{44} \end{aligned}$

Meta-analytic structural equation modeling

Combining meta-analyse (MA) and structural equation modeling (SEM)

Complete theoretical models Mediating variables Model fit Latent variables



Standard meta-analysis







Example

Eltanamly et al. (2021)



War \rightarrow Parenting behavior \rightarrow Child behavioral problems

Study 1



Study 2



Study k



Eltanamly et al. (2021) 38 studies





Methods for MASEM

Option 1: Pool correlations (or covariances), then fit SEM

- Univariate-r (Viswesvaran and Ones, 1995)
- GLS (Becker, 1992)
- Two Stage SEM (Cheung and Chan, 2005; Cheung, 2015)
- One-stage MASEM (Jak and Cheung, 2020)

Option 2: Fit SEM, then pool the SEM parameters

Becker and Wu (2007)

Gnambs and Staufenbiel (2016)

Disadvantage: Need complete data, and model should fit equally well in all samples

Ke, Zhang and Tong (2019)

Bayesian method, solves the issues with parameter-based MASEM

Methods for MASEM



Univariate-r approach

Stage 1: Use several univariate meta-analyses to pool each correlation coefficient



Univariate-r approach

Stage 2: Fit the path model on the pooled correlation matrix

	V1	V2	V3
V2	r ₂₁		
V3	r ₃₁	r ₃₂	
V4	r ₄₁	r ₄₂	r ₃₄



Univariate-r approach

Stage 1: Use several univariate meta-analyses to pool each correlation coefficient

- Ignores sampling covariance
- Can lead to non-positive definite correlation matrices

Stage 2: Fit the path model on the pooled correlation matrix using standard SEM software

- What is the sample size?
- Differences in precision of estimated correlations ignored
- Between-studies variance not taken into account

Multivariate approaches

GLS, TSSEM and one-stage MASEM involve multivariate metaanalysis of correlation coefficients



	V1	V2	V3
V2	r ₂₁		
V3	r ₃₁	r ₃₂	
V4	r ₄₁	r ₄₂	r ₃₄

Estimates of average correlations: $\hat{\rho}_R$ And between-study (co)variances of correlations across studies: \widehat{T}^2

Between-studies heterogeneity

GLS, TSSEM and one-stage MASEM

4 variables 6 mean correlations:

	V1	V2	V3
V2	r ₂₁		
V3	r ₃₁	r ₃₂	
V4	r ₄₁	r ₄₂	r ₄₃

Between-studies heterogeneity

GLS, TSSEM and one-stage MASEM

With 6 mean correlations:

Between-studies covariance matrix T²

6 variances 15 covariances

In practice: Use diagonal **T**² (Becker & Aloe, 2019)



TSSEM

Stage 1: Random effects multivariate meta-analysis of correlation coefficients using ML estimation

$\boldsymbol{r}_i = \boldsymbol{\rho}_R + \boldsymbol{u}_i + \boldsymbol{\varepsilon}_i$

- $cov(\boldsymbol{\varepsilon}_i) = \mathbf{V}_i$ (Within studies covariances)
- $cov(\mathbf{u}_i) = \mathbf{T}^2$ (Between studies (co)variances)

Stage 2: Fit SEM with WLS

$\psi_{11} \\ \psi_{21}$	Ψ22		
$\beta_{31}\psi_{11}\!+\!\beta_{32}\psi_{21}$	$\beta_{31}\psi_{21}\!+\!\beta_{32}\psi_{22}$	$ \begin{aligned} & \left(\beta_{31}\psi_{11}+\beta_{32}\psi_{21}\right)\beta_{31}+ \\ & \left(\beta_{31}\psi_{21}+\beta_{32}\psi_{22}\right)\beta_{32}+\psi_{33} \end{aligned} $	
$\begin{array}{c} \beta_{31}\beta_{43}\psi_{11}\texttt{+}\\ \beta_{32}\beta_{43}\psi_{21} \end{array}$	$\begin{array}{l} \beta_{31}\beta_{43}\psi_{21}\textbf{+}\\ \beta_{32}\beta_{43}\psi_{22} \end{array}$	$\begin{split} & (\beta_{31}\beta_{43}\psi_{11}\!+\!\beta_{32}\beta_{43}\psi_{21})\beta_{31}\!+\!\\ & (\beta_{31}\beta_{43}\psi_{21}\!+\!\beta_{32}\beta_{43}\psi_{22})\beta_{32}\!+\!\\ & \beta_{43}\psi_{33} \end{split}$	$\begin{split} &(\beta_{31}\beta_{43}\psi_{11}+\beta_{32}\beta_{43}\psi_{21})\beta_{31}\beta_{43}+\\ &(\beta_{31}\beta_{43}\psi_{21}+\beta_{32}\beta_{43}\psi_{22})\beta_{32}\beta_{43}+\\ &\beta_{43}{}^2\psi_{33}+\psi_{44} \end{split}$

$$\widehat{\boldsymbol{F}}_{\text{WLS}} = (\widehat{\boldsymbol{\rho}}_R - \widehat{\boldsymbol{\rho}}_{MODEL})^{\mathsf{T}} \widehat{\boldsymbol{V}}_R^{-1} (\widehat{\boldsymbol{\rho}}_R - \widehat{\boldsymbol{\rho}}_{MODEL})$$

	V1	V2	V3	V4
V1	σ_{11}			
V2	σ_{21}	σ_{22}		
V3	σ_{31}	σ_{32}	σ_{33}	
V4	σ_{41}	σ_{42}	σ_{43}	σ_{44}

GLS approach

Stage 1: Random effects multivariate meta-analysis of correlation coefficients

$$r = \rho_R + u + \varepsilon$$

$$\widehat{\boldsymbol{\rho}}_R = (\mathsf{X}^{\top} \, \mathsf{\Sigma}^{-1} \, \mathsf{X})^{-1} \, \mathsf{X}^{\top} \, \mathsf{\Sigma}^{-1} \, \mathsf{r}$$
 $\widehat{\boldsymbol{V}}_R = (\mathsf{X}^{\top} \, \mathsf{\Sigma}^{-1} \mathsf{X})^{-1}$

r = vector of observed correlations **X** = stacked selection matrices **Σ** = block-diagonal matrix with $V_i + T^2$

Stage 2: $\widehat{B} = \widehat{\rho}_{XX} - \widehat{\rho}_{YX}$

Alternative: use WLS with $\widehat{oldsymbol{
ho}}_R$ and $\widehat{oldsymbol{V}}_R$ like in TSSEM

One-stage MASEM



One-stage MASEM

Multivariate random-effects analysis of correlation coefficients

$$r_i = \rho_R + u_i + \varepsilon_i$$

One-stage MASEM restricts the pooled correlation matrix to a SEM model:

$$\boldsymbol{\rho}_R = \operatorname{vechs}(\boldsymbol{\Lambda}\boldsymbol{\Phi}\boldsymbol{\Lambda}^{-1t} + \boldsymbol{\Theta})$$

The SEM parameters may be regressed on study-level moderator variables

Comparison of four methods

Data generating model

Based on Nohe et al. (2015)



Fit correct model using univariate-r, GLS, TSSEM and one-stage MASEM

Comparison of four methods

Fit correct path model using

Univariate-r

Univariate meta-analyses of raw correlations, weighted by N Using harmonic mean as sample size in Stage 2

Multivariate methods (GLS, TSSEM, one-stage MASEM) With diagonal T² (although generated full T²)

Conditions

Varying:

- Number of studies
 - k = 16, 32 or 64
- Missing data

0, 50 or 75% of the studies missed 2 of the 4 variables

- Heterogeneity (**T**²)

Between-studies SD .05 or .10 Between studies correlations .10 or .30

Fixed:

- Sample size per study

Evaluation

Rejection percentages of test statistic Relative bias in parameter estimates Relative bias in standard errors

Test statistic (rejection percentages)

	- i	Pop	oulation i	rho SD=	=0.05	Population rho SD=0.05				Po	pulation	rho SD:	=0.1	Population rho SD=0.1					
	Populaton rho cor=0.1				Populaton rho cor=0.3			Po	Populaton rho cor=0.1			Populaton rho cor=0.3							
	64 -	4.90	5.70	6.60	49.35	2.70	4.70	4.90	45.29	4.20	4.50	4.60	79.50	2.20	3.60	3.80	77.56	2	
	32 -	5.01	5.40	6.30	49.30	3.00	4.60	5.22	43.90	3.70	3.80	4.60	83.70	2.50	3.80	4.30	78.90	lissing studies:	
	16 -	6.26	6.50	8.02	44.74	5.15	5.50	6.89	39.00	6.50	6.70	7.40	77.30	2.50	3.80	5.30	76.10	=0	
s (k)	64 -	4.30	4.70	5.73	49.70	4.21	6.10	6.11	41.50	4.20	4.80	5.40	80.50	3.00	4.00	4.10	77.58	M	Rejection rate
per of studie:	32 -	6.71	6.40	7.54	46.40	3.50	5.20	5.34	39.00	5.00	5.10	6.01	79.00	2.50	3.60	4.40	72.40	ssing studies=	9 8 7 6 5 4 3
Num	16 -	8.19	7.60	9.06	43.80	5.47	5.90	5.99	35.90	6.50	5.60	8.66	74.70	5.11	6.20	7.72	73.90	0.5	2 1 0
					-				-				-				-		
	64 -	6.60	5.90	6.83	43.90	4.10	5.20	5.93	41.80	4.60	5.00	6.01	78.50	2.80	3.50	3.80	73.40	Mi	
	32 -	7.41	6.50	6.39	43.20	5.32	5.90	6.17	39.00	6.60	6.30	9.01	75.10	5.00	6.60	7.53	76.00	ssing studies=	
	16 -	10.52	8.90	9.24	35.60	9.74	9.00	8.93	34.30	8.50	7.00	12.68	70.90	8.00	7.10	10.56	66.80	-0.75	
		gls o	smaser	ntssem	unir	gls (osmaser	ntssem	unir	gls (osmaser	ntssem	unir	gls	osmase	mtssem	unir		

Rejection rates of the test statistics (expected counts=5 for alpha=.05)

Method

Bias in parameter estimates

K = 32, SD = 0.1, cor = 0.3, missing studies = 0.5



Bias in parameter estimates

Relative percentage bias of parameter estimates



Bias in standard errors

K = 32, SD = 0.1, cor = 0.3, missing studies = 0.5



Bias in standard errors

Relative percentage bias of standard errors



Conclusions

Univariate-r approach leads to unbiased parameter 🗸 estimates, but:

Extremely inflated test-statistics and associated Type 1 🗙 errors

Extremely biased standard errors



The three multivariate methods generally lead to unbiased parameter estimates, well behaved test-statistics, and correct standard errors



Within the Journal of Applied Psychology, all-but-one of the MASEM applications since 2020 used the univariate-r method

Multivariate methods also need more research

Minimum sample size conditions (e.g., how many studies are needed in which conditions?)

Handling dependent effect sizes

However, it is worrying how often researchers apply the only method that clearly leads to wrong results in all situations

Why is the univariate method (still) so popular?

Because it is easier to apply?

Not anymore!

Tutorial and Shiny app for one-stage MASEM (Jak et al. 2021)

https://sjak.shinyapps.io/webMASEM/



Why is the univariate method (still) so popular?

Because researchers copy the current practice I their field

Harder to change, but we are hopeful.

Thanks for listening!

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