

The ‘Structural After Measurement’ (SAM) approach to SEM

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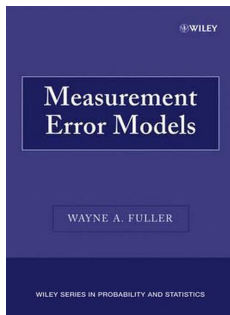
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acknowledgements

- Dr. Wen Wei Loh (Maastricht University)
- Dr. Sara Dhaene (UGent)

inspiration

Fuller, W.A. (1987). *Measurement Error Models*. Wiley.



the SAM paper

Rosseel, Y., & Loh, W.W. (2022). A structural after measurement approach to structural equation modeling. *Psychological Methods*

<https://doi.org/10.1037/met0000503>

<https://osf.io/pekbm/> (includes original version)

related literature

Levy, R. (2023). Precluding interpretational confounding in factor analysis with a covariate or outcome via measurement and uncertainty preserving parametric modeling. *Structural Equation Modeling: A Multidisciplinary Journal*.

<https://doi.org/10.1080/10705511.2022.2154214>

Bakk, Z., & Kuha, J. (2021). Relating latent class membership to external variables: An overview. *British Journal of Mathematical and Statistical Psychology*, 74(2), 340–362.

<https://doi.org/10.1111/bmsp.12227> (Open Access)

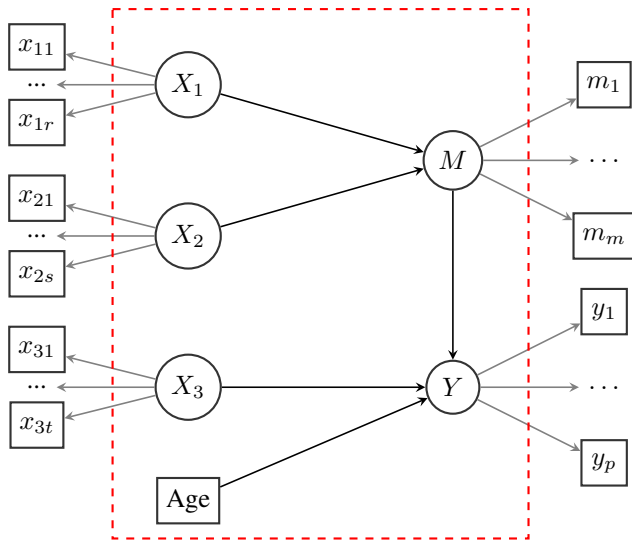
Kuha, J., & Bakk, Z. (arxiv.org). Two-step estimation of latent trait models.

<https://arxiv.org/pdf/2303.16101.pdf>

the setting

- structural equation models
- we focus on ‘large’ models with many (say, > 100) parameters:
 - many constructs (motivation, ability, personality traits, ...)
 - each construct is measured by a set of (observed) indicators
 - many ‘background’ variables (age, gender, ...)
 - multilevel data, missing data, ...
- we are mostly interested in the structural part of the model:
 - regression model: variables are either dependent or independent
 - path analysis model: includes mediating effects, perhaps non-recursive
- assumption: the measurement instruments for the latent variables are well established, and usually fit (reasonably) well
- BUT: the sample size is not large (say, $N = 150$)

the setting

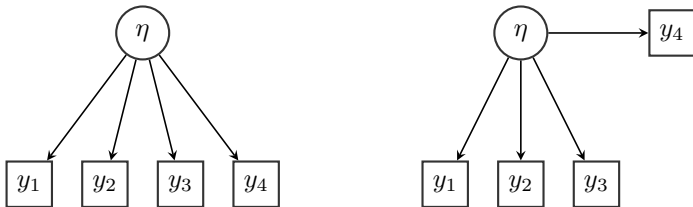


model parameters

- the majority of the model parameters are related to the measurement part of the model:
 - factor loadings
 - residual variances for the indicators
 - factor (co)variances
- a small portion of the model parameters are related to the structural part of the model:
 - regression coefficients
 - (residual) (co)variances

the standard estimation approach in SEM

- all parameters (measurement and structural) are estimated jointly: “system-wide” estimation
- frequentist: typically using an iterative optimization approach (e.g., ML)
- Bayesian: typically using MCMC
- advantages:
 - one-step
 - efficient (in terms of sampling variability)
 - inference is straightforward (standard errors, hypothesis testing)
 - (relatively) easy to handle constraints, missing data, . . .
- works very well if the following conditions are met:
 - correctly specified model
 - large sample size
 - (normally distributed data)

example (stolen from Roy Levy, 2023)

- left panel: only measurement
- right panel: measurement + structural
- mathematically identical (in standard SEM)
- conceptually very different

generate some data

```
> library(lavaan)
> Sigma <- matrix(c(2.0, 1.0, 1.0, 1.0, 1.5,
                  1.0, 2.0, 1.0, 1.0, 0.1,
                  1.0, 1.0, 2.0, 1.0, 0.1,
                  1.0, 1.0, 1.0, 2.0, 0.1,
                  1.5, 0.1, 0.1, 0.1, 2.0), nrow = 5, ncol = 5)
> rownames(Sigma) <- colnames(Sigma) <- c("y1", "y2", "y3", "y4", "z")
> Sigma
```

```
      y1 y2 y3 y4 z
y1 2.0 1.0 1.0 1.0 1.5
y2 1.0 2.0 1.0 1.0 0.1
y3 1.0 1.0 2.0 1.0 0.1
y4 1.0 1.0 1.0 2.0 0.1
z   1.5 0.1 0.1 0.1 2.0
```

```
> set.seed(3)
> Data <- MASS::mvrnorm(n = 200L, mu = rep(0, 5), Sigma = Sigma)
```

R code left panel (model1)

```
> model1 <- '
  f =~ y1 + y2 + y3 + y4
,
> fit1 <- sem(model1, data = Data)
> summary(fit1)
```

lavaan 0.6.18.2004 ended normally after 25 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	8
Number of observations	200

Model Test User Model:

Test statistic	0.140
Degrees of freedom	2
P-value (Chi-square)	0.932

Parameter Estimates:

Standard errors	Standard
Information	Expected
Information saturated (h1) model	Structured

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
f =~				
y1	1.000			
y2	0.996	0.123	8.122	0.000
y3	0.894	0.116	7.689	0.000
y4	0.911	0.120	7.608	0.000

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y1	1.096	0.148	7.388	0.000
.y2	0.837	0.127	6.601	0.000
.y3	0.987	0.128	7.694	0.000
.y4	1.081	0.138	7.820	0.000
f	1.031	0.208	4.959	0.000

R code right panel (model2)

```
> model2 <- '
  f =~ y1 + y2 + y3
  y4 ~ f
',
> fit2 <- sem(model2, data = Data)
> summary(fit2)
```

lavaan 0.6.18.2004 ended normally after 26 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	8
Number of observations	200

Model Test User Model:

Test statistic	0.140
Degrees of freedom	2
P-value (Chi-square)	0.932

Parameter Estimates:

Standard errors	Standard
Information	Expected
Information saturated (h1) model	Structured

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
f =~				
y1	1.000			
y2	0.996	0.123	8.122	0.000
y3	0.894	0.116	7.689	0.000

Regressions:

	Estimate	Std.Err	z-value	P(> z)
y4 ~				
f	0.911	0.120	7.608	0.000

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y1	1.096	0.148	7.388	0.000
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.y3	0.987	0.128	7.694	0.000
.y4	1.081	0.138	7.820	0.000
f	1.031	0.208	4.959	0.000

change outcome variable (y4 becomes z) (model3)

```
> model3 <- '
  f =~ y1 + y2 + y3
  z =~ f
'
> fit3 <- sem(model3, data = Data)
> summary(fit3)
```

lavaan 0.6.18.2004 ended normally after 58 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	8
Number of observations	200

Model Test User Model:

Test statistic	50.659
Degrees of freedom	2
P-value (Chi-square)	0.000

Parameter Estimates:

Standard errors	Standard
Information	Expected
Information saturated (h1) model	Structured

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
f =~				
y1	1.000			
y2	0.138	0.053	2.601	0.009
y3	0.133	0.051	2.591	0.010

Regressions:

	Estimate	Std.Err	z-value	P(> z)
z ~				
f	0.276	0.087	3.156	0.002

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y1	-4.169	1.801	-2.315	0.021
.y2	1.740	0.173	10.071	0.000
.y3	1.699	0.168	10.086	0.000
.z	1.677	0.211	7.953	0.000
f	6.296	1.715	3.671	0.000

interpretational confounding

- replacing y_4 by z (= changing the structural part) also changes the parameters of the measurement model
- if the resulting parameters of the measurement model imply a different ‘meaning’ of the latent variable than was intended by the researcher, we have a problem
- this problem was coined “interpretational confounding” by Burt (1976)

Burt, R.S. (1976). Interpretational confounding of unobserved variables in structural equation models. *Sociological Methods & Research*, 5(1), 3–52.

- Burt (1976) already suggested the solution: first fit the measurement part of the model, and then fit the structural part of the model

solution: replace sem() by sam()

```
> fit3.sam <- sam(model3, data = Data)
> parameterEstimates(fit3.sam, remove.step1 = FALSE, ci = FALSE,
  output = "text")
```

Latent Variables:

	Step	Estimate	Std.Err	z-value	P(> z)
f =~					
y1	1	1.000			
y2	1	0.969	0.139	6.977	0.000
y3	1	0.884	0.127	6.979	0.000

Regressions:

	Step	Estimate	Std.Err	z-value	P(> z)
z ~					
f	2	0.540	0.121	4.471	0.000

Variances:

	Step	Estimate	Std.Err	z-value	P(> z)
.y1	1	1.067	0.170	6.287	0.000
.y2	1	0.865	0.151	5.723	0.000
.y3	1	0.983	0.142	6.900	0.000
.z	2	1.846	0.195	9.462	0.000
f	2	1.059	0.226	4.678	0.000

the structural-after-measurement (SAM) approach

- SAM is an umbrella term to describe many different (estimation) approaches that have the following in common:
 - in the first step: we estimate the parameters related to the measurement part (factor loadings, residual variance of the indicators)
 - in the second step: we estimate the parameters related to the structural part (regression coefficients, residual (co)variances)
- the term SAM was used by Rosseel & Loh (2022), to avoid the overloaded terms ‘two-step’, ‘two-stage’, ...
- various SAM approaches have been suggested in the literature:
 - early references: Burt (1976), Hunter & Gerbing (1982), Lance, Cornwell & Mulaik (1988)
 - (uncorrected and bias-corrected) factor score regression (FSR)
 - SAM is the default approach in many other fields
- ... but they never received much attention in the SEM literature/community

critique on the SAM approach

- the (naive) standard errors in the second step are wrong (because they ignore the uncertainty that stems from the first step)
- in general: inference is (more) complicated
- multiple step methods are less efficient (more sampling variability)
- Fornell and Yi (1992) gave an example where a misspecified (but well-fitting) measurement model was embedded in a correctly specified structural model; but the model fit of the full model suggested that the model did not fit well, thus incorrectly implying a misspecified structural model
- software packages only allow for joint estimation
- ...
- to address (most of) these issues, we developed the `sam()` function in `lavaan`
- per default, `sam()` uses a method called 'local' SAM

local SAM: rationale

- the measurement model:

$$\mathbf{y} = \boldsymbol{\nu} + \boldsymbol{\Lambda}\boldsymbol{\eta} + \boldsymbol{\epsilon}$$

- to solve this for $\boldsymbol{\eta}$, we proceed as follows:

$$\boldsymbol{\nu} + \boldsymbol{\Lambda}\boldsymbol{\eta} + \boldsymbol{\epsilon} = \mathbf{y}$$

$$\boldsymbol{\Lambda}\boldsymbol{\eta} = \mathbf{y} - \boldsymbol{\nu} - \boldsymbol{\epsilon}$$

$$\mathbf{M}\boldsymbol{\Lambda}\boldsymbol{\eta} = \mathbf{M}[\mathbf{y} - \boldsymbol{\nu} - \boldsymbol{\epsilon}]$$

$$\boldsymbol{\eta} = \mathbf{M}[\mathbf{y} - \boldsymbol{\nu} - \boldsymbol{\epsilon}]$$

where \mathbf{M} is $M \times P$ mapping matrix such that $\mathbf{M}\boldsymbol{\Lambda} = \mathbf{I}_M$

- we assume $\mathbf{E}(\boldsymbol{\epsilon}) = \mathbf{0}$ and write $\text{Var}(\boldsymbol{\epsilon}) = \boldsymbol{\Theta}$; it follows that

$$\mathbf{E}(\boldsymbol{\eta}) = \mathbf{M}[\mathbf{E}(\mathbf{y}) - \boldsymbol{\nu}]$$

$$\text{Var}(\boldsymbol{\eta}) = \mathbf{M}[\text{Var}(\mathbf{y}) - \boldsymbol{\Theta}] \mathbf{M}^T$$

local SAM: first stage

- first stage: estimation of the measurement part of the model (only)
- this results in estimates for ν , Λ and Θ
- M is the number of latent variables; B is the number of measurement ‘blocks’
- three options:
 1. $B = 1$: single CFA
 2. $B = M$: as many ‘blocks’ as we have latent variables
 3. $B < M$: if some blocks are ‘linked’ together
- measurement models that are ‘linked’ (due to cross-loadings, correlated residuals, or equality constraints) should be fitted together
- measurement models that are ‘weak’ (low construct reliability) should also be fitted together in order to ‘borrow strength’ from each other
- for each block, we can use ML, GLS, ..., or we can use noniterative estimators

local SAM: creating the mapping matrix \mathbf{M}

- recall, the mapping matrix must be chosen such that $\mathbf{M}\mathbf{\Lambda} = \mathbf{I}_M$
- three possible solutions for the mapping matrix \mathbf{M} :

$$\mathbf{M} = (\mathbf{\Lambda}^T \mathbf{\Theta}^{-1} \mathbf{\Lambda})^{-1} \mathbf{\Lambda}^T \mathbf{\Theta}^{-1} \quad (ML)$$

$$\mathbf{M} = (\mathbf{\Lambda}^T \mathbf{S}^{-1} \mathbf{\Lambda})^{-1} \mathbf{\Lambda}^T \mathbf{S}^{-1} \quad (GLS)$$

$$\mathbf{M} = (\mathbf{\Lambda}^T \mathbf{\Lambda})^{-1} \mathbf{\Lambda}^T \quad (ULS)$$

- we then estimate $E(\boldsymbol{\eta})$ and $\text{Var}(\boldsymbol{\eta})$ as follows:

$$\widehat{E}(\boldsymbol{\eta}) = \hat{\mathbf{M}} [\bar{\mathbf{y}} - \hat{\boldsymbol{\nu}}]$$

$$\widehat{\text{Var}}(\boldsymbol{\eta}) = \hat{\mathbf{M}} [\mathbf{S} - \hat{\boldsymbol{\Theta}}] \hat{\mathbf{M}}^T$$

- in local SAM, we proceed in the second stage with these ‘sufficient statistics’ only

local SAM: second stage

- second stage: $\widehat{E}(\boldsymbol{\eta})$ and $\widehat{\text{Var}}(\boldsymbol{\eta})$ are used to estimate structural parameters of the model
- this can be done using ‘path analysis’, where we treat everything as observed, and the data is presented via summary statistics
- we can use ML, GLS, ...
- or we can use noniterative estimators: OLS (if the model is recursive) or TSLS (if the model is not recursive)

local SAM: further comments

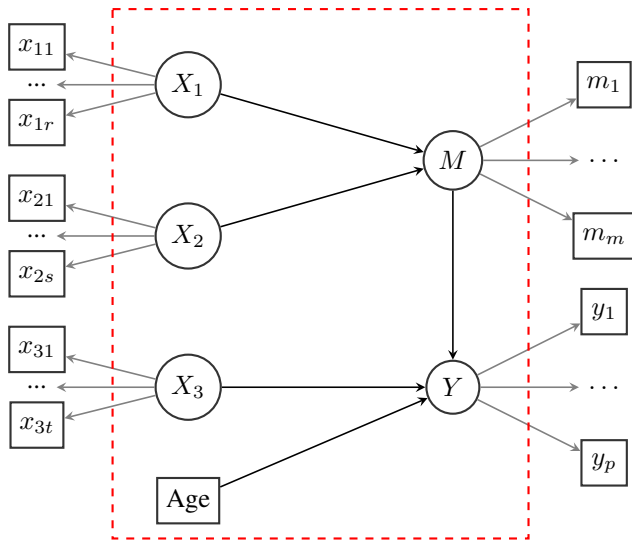
- two-step corrected standard errors are available (seen Appendix C in the SAM paper)
- local fit measures only (for each measurement block, for the structural part)
- the (co)variance matrix of the latent variables is always positive definite
- we can handle missing data (fiml or two-stage), categorical indicators, twolevel data (random intercepts only), ...
- but still some limitations (we plan to address these limitations in future work):
 - all indicators (of latent constructs) must be observed (i.e., no second-order measurement models)
 - the factor loading matrix (Λ) must have full column rank
 - no support for (e.g., variance components) models where zeroes in the variance-covariance matrix of the latent variables are needed in order to identify the model

example: generate population data mediation model

```
> library(lavaan)
> pop.model <- '
  # factor loadings
  Y  =~ 1*y1 + 1.2*y2 + 0.8*y3 + 0.5*y4
  M  =~ 1*m1 + 0.5*m2 + 0.5*m3 + 0.7*m4
  X1 =~ 1*x1 + 0.7*x2 + 0.6*x3 + 1.1*x4
  X2 =~ 1*x5 + 0.7*x6 + 0.6*x7 + 0.9*x8
  X3 =~ 1*x9 + 0.7*x10 + 0.6*x11 + 1.1*x12

  # covariances among exogenous X1-X3
  X1 ~~ 0.4*X2; X1 ~~ -0.2*X3; X2 ~~ 0.4*X3

  # regression part
  Y  ~ 0.25*X3 + 0.4*M + (-0.1)*Age
  M  ~ -0.30*X1 + 1.1*X2
  ,
> set.seed(1234)
> Data <- simulateData(pop.model, sample.nobs = 200L, empirical = TRUE)
```

example: diagram

example: fitting the model using traditional SEM

```

> model <- '
  # measurement part
  Y =~ y1 + y2 + y3 + y4
  M =~ m1 + m2 + m3 + m4
  X1 =~ x1 + x2 + x3 + x4
  X2 =~ x5 + x6 + x7 + x8
  X3 =~ x9 + x10 + x11 + x12

  # structural part
  Y ~ X3 + M + Age
  M ~ X1 + X2
,
> fit.sem <- sem(model, data = Data, estimator = "ML")
> parameterEstimates(fit.sem, ci = FALSE, output = "text")[21:25,]

```

Regressions:

	Estimate	Std.Err	z-value	P(> z)
Y ~				
X3	0.250	0.108	2.308	0.021
M	0.400	0.079	5.078	0.000
Age	-0.100	0.083	-1.203	0.229
M ~				
X1	-0.300	0.133	-2.258	0.024
X2	1.100	0.165	6.670	0.000

example: model-implied variance-covariance matrix latent variables

```
> lavInspect (fit.sem, "cov.lv")
```

	Y	M	X1	X2	X3
Y	1.498				
M	0.939	2.036			
X1	0.006	0.140	1.000		
X2	0.492	0.980	0.400	1.000	
X3	0.450	0.500	-0.200	0.400	1.000

example: fit measurement blocks, using $B = M$

```
> fit.Y <- sem('Y =~ y1 + y2 + y3 + y4', data = Data)
> fit.M <- sem('M =~ m1 + m2 + m3 + m4', data = Data)
> fit.X1 <- sem('X1 =~ x1 + x2 + x3 + x4', data = Data)
> fit.X2 <- sem('X2 =~ x5 + x6 + x7 + x8', data = Data)
> fit.X3 <- sem('X3 =~ x9 + x10 + x11 + x12', data = Data)

> # assemble Lambda and Theta
> Lambda <- matrix(0, 20, 5)
> Lambda[ 1:4, 1] <- lavInspect(fit.Y, "est")$lambda
> Lambda[ 5:8, 2] <- lavInspect(fit.M, "est")$lambda
> Lambda[ 9:12, 3] <- lavInspect(fit.X1, "est")$lambda
> Lambda[13:16, 4] <- lavInspect(fit.X2, "est")$lambda
> Lambda[17:20, 5] <- lavInspect(fit.X3, "est")$lambda

> Theta <- lav_matrix_bdiag(lavInspect(fit.Y, "est")$theta,
                           lavInspect(fit.M, "est")$theta,
                           lavInspect(fit.X1, "est")$theta,
                           lavInspect(fit.X2, "est")$theta,
                           lavInspect(fit.X3, "est")$theta)
```

example: compute ML version of the mapping matrix M

```

> Theta.inv <- solve(Theta)
> M <- solve(t(Lambda) %**% Theta.inv %**% Lambda) %**% t(Lambda) %**% Theta.inv

> # add age
> M      <- lav_matrix_bdiag(M,      matrix(1, nrow = 1L, ncol = 1L))
> Theta <- lav_matrix_bdiag(Theta, matrix(0, nrow = 1L, ncol = 1L))
> rownames(M) <- c("Y", "M", "X1", "X2", "X3", "Age")

> # compute (biased) sample covariance matrix 'S'
> N <- nrow(Data)
> S <- cov(Data) * (N - 1L)/N

> # compute Var(Eta)
> Var.eta <- M %**% (S - Theta) %**% t(M)
> round(Var.eta, 3)

```

	Y	M	X1	X2	X3	Age
Y	1.498	0.939	0.006	0.492	0.45	-0.1
M	0.939	2.036	0.140	0.980	0.50	0.0
X1	0.006	0.140	1.000	0.400	-0.20	0.0
X2	0.492	0.980	0.400	1.000	0.40	0.0
X3	0.450	0.500	-0.200	0.400	1.00	0.0
Age	-0.100	0.000	0.000	0.000	0.00	1.0

example: second stage – using OLS

```
> # compute regression coefficients for M
> beta.M <- ( solve(Var.eta[c("X1", "X2"), c("X1", "X2")]) %*%
              Var.eta[c("X1", "X2"), "M", drop = FALSE] )
> round(beta.M, 3)
```

```
      M
X1 -0.3
X2  1.1
```

```
> # compute regression coefficients for Y
> beta.Y <- ( solve(Var.eta[c("X3", "M", "Age"), c("X3", "M", "Age")]) %*%
              Var.eta[c("X3", "M", "Age"), "Y", drop = FALSE] )
> round(beta.Y, 3)
```

```
      Y
X3  0.25
M   0.40
Age -0.10
```

example: using the sam() function

```
> fit.lsam <- sam(model = model, data = Data)
> parameterEstimates(fit.lsam, ci = FALSE, output = "text")[1:5,]
```

Regressions:

	Estimate	Std.Err	z-value	P(> z)
Y ~				
X3	0.250	0.109	2.301	0.021
M	0.400	0.080	4.971	0.000
Age	-0.100	0.083	-1.203	0.229
M ~				
X1	-0.300	0.133	-2.251	0.024
X2	1.100	0.176	6.235	0.000

application 1: adding latent quadratic and interaction terms

- in the joint setting, adding latent quadratic/interaction terms is not trivial
- two popular methods are the product-indicator (PI) approach, and the so-called 'Latent Moderated Structural Equations' (LMS) approach
- none of these scale well: they cannot handle many quadratic and latent interaction terms simultaneously
- but if you can decouple the measurement and structural part, this becomes feasible
- a very general SAM solution (allowing for polynomial relations between latent variables) was already described in Wall & Amemiya (2000)
- the local SAM approach: find an explicit expression for

$$E(\boldsymbol{\eta} \otimes \boldsymbol{\eta}) \quad \text{and} \quad \text{Var}(\boldsymbol{\eta} \otimes \boldsymbol{\eta})$$

where \otimes denotes the tensor (or Kronecker) product

implementation in lavaan

```
> model <- '  
  # measurement part  
  f1 =~ y1 + y2 + y3  
  f2 =~ y4 + y5 + y6  
  f3 =~ y7 + y8 + y9  
  
  # structural part  
  f3 ~ f1 + f2 + f1:f1 + f2:f2 + f1:f2  
,  
> fit <- sam(model, data = Data, se = "none") # or se = "naive"
```

- no two-step analytic standard errors yet; but bootstrapping is possible
- forthcoming paper:

Rosseel, Y., Burghraeve, E., Loh, W.W., Schermelleh-Engel, K. (accepted). Structural after Measurement (SAM) approaches for accommodating latent quadratic and interaction effects. *Behavior Research Methods*.

application 2: noniterative SEM

- for CFA, many noniterative estimators are available; some (i.e., the multiple group method) perform better than ML in terms of mean squared error

Dhaene, S. & Rosseel, Y. (2023). An Evaluation of Non-Iterative Estimators in Confirmatory Factor Analysis. *Structural Equation Modeling: A Multidisciplinary Journal*.

- we can use these noniterative estimators for the measurement part in SAM

Dhaene, S., & Rosseel, Y. (2023). An Evaluation of Non-Iterative Estimators in the Structural after Measurement (SAM) Approach to Structural Equation Modeling (SEM). *Structural Equation Modeling: A Multidisciplinary Journal*, 30(6), 926–940

- “[the] local SAM approach outperforms traditional SEM in small to moderate samples (both in terms of convergence and MSE values), especially when reliability drops. ”
- BUT: no analytic standard errors yet (pseudo-ML or bootstrapping)

application 3: comparing structural relations across many groups

- reference:

Perez Alonso, A.F., Rosseel, Y., Vermunt, J.K., & De Roover, K. (in press). Mixture Multigroup Structural Equation Modeling: A Novel Method for Comparing Structural Relations Across Many Groups. *Psychological Methods*.

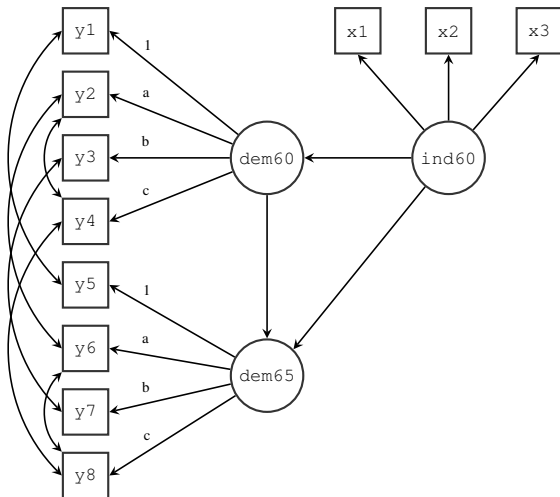
- relationships between latent variables are often different across groups (e.g., countries); but some groups may be similar in the sense that they have similar values for the regression coefficients
- we like to ‘discover’ these hidden clusters of similar groups
- in a first step, we estimated the measurement part across all groups (fixing the factor loadings to be the same across groups); this resulted in (model-implied) latent (co)variance matrices for all the groups
- in a second step, a mixture modeling approach is used to find homogeneous clusters that share similar regression coefficients

SAM: software implementation

- the SAM approach has been implemented in the `sam()` function in the R package `lavaan`
- available methods:
 - `sam.method = "local"` (default)
 - `sam.method = "global"`
 - `sam.method = "fsr"` (using Bartlett factor scores)
- typical call:

```
> fit.sam <- sam(model, data = PoliticalDemocracy,
  sam.method = "local",
  # link measurement blocks
  mm.list = list(ind = "ind60", dem = c("dem60", "dem65")),
  # measurement options
  mm.args = list(estimator = "ML"),
  # structural options
  struc.args = list(estimator = "GLS"),
  # global options
  meanstructure = FALSE)
```

diagram of the Political Democracy model



sam() output

```
> summary(fit.sam, remove.step1 = FALSE)
```

This is lavaan 0.6.18.2004 -- using the SAM approach to SEM

SAM method	LOCAL
Mapping matrix M method	ML
Number of measurement blocks	2
Estimator measurement part	ML
Estimator structural part	GLS
Number of observations	75

Summary Information Measurement + Structural:

Block	Latent	Nind	Chisq	Df
1	ind60	3	0.00	0
2	dem60, dem65	8	15.32	16

Model-based reliability latent variables:

ind60	dem60	dem65
0.966	0.868	0.87

Summary Information Structural part:

chisq	df	cfi	rmsea	srmr
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0 0 1 0 0

Parameter Estimates:

Standard errors Information Information saturated (h1) model	Twostep Expected Structured
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Latent Variables:

	Step	Estimate	Std.Err	z-value	P(> z)
ind60 =~					
x1	1	1.000			
x2	1	2.193	0.142	15.403	0.000
x3	1	1.824	0.153	11.883	0.000
dem60 =~					
y1	1	1.000			
y2 (a)	1	1.213	0.143	8.483	0.000
y3 (b)	1	1.210	0.125	9.690	0.000
y4 (c)	1	1.273	0.122	10.453	0.000
dem65 =~					
y5	1	1.000			
y6 (a)	1	1.213	0.143	8.483	0.000
y7 (b)	1	1.210	0.125	9.690	0.000
y8 (c)	1	1.273	0.122	10.453	0.000

Regressions:

	Step	Estimate	Std.Err	z-value	P(> z)
dem60 ~					

ind60	2	1.454	0.389	3.741	0.000
dem65 ~					
ind60	2	0.558	0.225	2.480	0.013
dem60	2	0.871	0.076	11.497	0.000

Covariances:

	Step	Estimate	Std.Err	z-value	P(> z)
.y1 ~~					
.y5	1	0.577	0.364	1.585	0.113
.y2 ~~					
.y4	1	1.390	0.685	2.030	0.042
.y6	1	2.068	0.733	2.822	0.005
.y3 ~~					
.y7	1	0.727	0.611	1.190	0.234
.y4 ~~					
.y8	1	0.476	0.453	1.049	0.294
.y6 ~~					
.y8	1	1.257	0.583	2.156	0.031

Variances:

	Step	Estimate	Std.Err	z-value	P(> z)
.x1	1	0.084	0.020	4.140	0.000
.x2	1	0.108	0.074	1.455	0.146
.x3	1	0.468	0.091	5.124	0.000
.y1	1	1.879	0.431	4.355	0.000
.y2	1	7.530	1.363	5.523	0.000
.y3	1	4.966	0.966	5.141	0.000
.y4	1	3.214	0.722	4.449	0.000

.y5	1	2.499	0.518	4.824	0.000
.y6	1	4.809	0.924	5.202	0.000
.y7	1	3.302	0.699	4.722	0.000
.y8	1	3.227	0.720	4.482	0.000
ind60	2	0.446	0.087	5.135	0.000
.dem60	2	3.766	0.848	4.439	0.000
.dem65	2	0.189	0.224	0.843	0.399

discussion

- why should we decouple the measurement and structural part?
 - because we should (avoid interpretational confounding)
 - because we can (we can still do SEM)
 - this is what is done in most fields outside SEM
 - good performance in simulation studies (small/moderate sample sizes)
 - opens up modeling possibilities that were (computationally) difficult (if not impossible) in a joint estimation approach
- but still some obstacles:
 - analytic standard errors not always available (yet)
 - limitations (e.g., no higher-order measurement models)
 - more study is needed to discover potential weaknesses
- the SAM approach deserves the (renewed) interest of the SEM community

Thank you!

(questions?)

<https://lavaan.org>

<https://lavaan.ugent.be/about/donate.html>

[https://jobs.ugent.be/job/
Ghent-Post-doctoral-researcher-9000/786525702/](https://jobs.ugent.be/job/Ghent-Post-doctoral-researcher-9000/786525702/)

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