

Structural Equation Modeling with lavaan

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Before we start ...

- slides (and all other files) are available here:

<https://users.ugent.be/~yrosseel/lavaan/maastricht2024>

- in the interest of time, we will only spend a short time for the ‘practical session’
- but after this course, you can use the `lavaan_labs.pdf` document to practice what you have learned
- a starting script (`lavaan_labs_start.R`) and a feedback script (`lavaan_labs_feedback.R`) are also available

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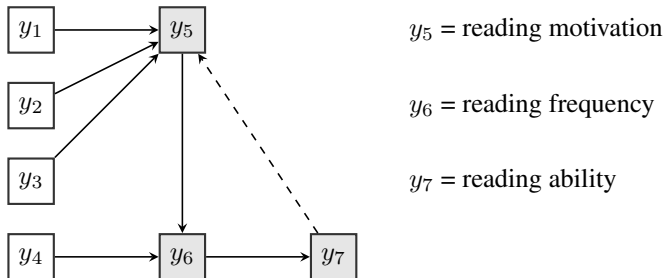
1 Introduction to SEM

1.1 What is SEM?

- SEM is a multivariate statistical modeling technique
- SEM allows us to test a hypothesis/model about the data
 - we postulate a data-generating model
 - this model may or may not fit the data
- what is so special about SEM?
 1. the model may contain latent variables
 - latent variables can be hypothetical ‘constructs’ (eg., depression) measured by a set of indicators
 - latent variables can be random effects (eg., random intercepts)
 - error terms, missing data, . . .
 2. SEM allows for indirect effects (mediation), reciprocal effects, . . .
 3. the model is depicted as a diagram

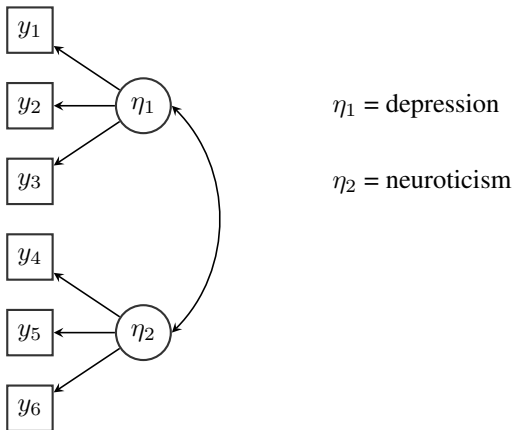
path analysis

- all variables are observed (manifest)
- we allow for indirect effects (eg., of y_5 , via y_6 on y_7)
- we allow for cycles (eg. y_7 could influence y_5)



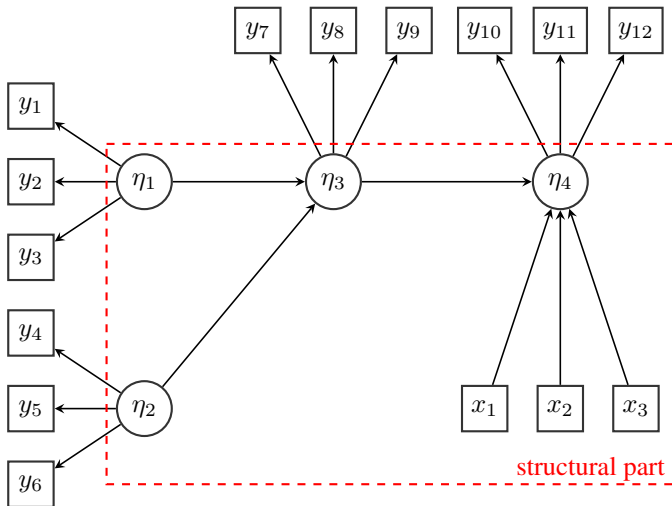
confirmatory factor analysis (CFA)

- measurement model: representing the relationship between one or more latent variables and their (observed) indicators



structural equation modeling (SEM)

- path analysis with latent variables



1.2 How does SEM work?

a dataset: the Holzinger & Swineford dataset

```
> library(lavaan)
> var.names <- c("x1", "x2", "x3", "x4", "x5", "x6", "x7", "x8", "x9")
> summary(HolzingerSwineford1939[, var.names])
```

x1		x2		x3		x4	
Min.	:0.6667	Min.	:2.250	Min.	:0.250	Min.	:0.000
1st Qu.	:4.1667	1st Qu.	:5.250	1st Qu.	:1.375	1st Qu.	:2.333
Median	:5.0000	Median	:6.000	Median	:2.125	Median	:3.000
Mean	:4.9358	Mean	:6.088	Mean	:2.250	Mean	:3.061
3rd Qu.	:5.6667	3rd Qu.	:6.750	3rd Qu.	:3.125	3rd Qu.	:3.667
Max.	:8.5000	Max.	:9.250	Max.	:4.500	Max.	:6.333
x5		x6		x7		x8	
Min.	:1.000	Min.	:0.1429	Min.	:1.304	Min.	:3.050
1st Qu.	:3.500	1st Qu.	:1.4286	1st Qu.	:3.478	1st Qu.	:4.850
Median	:4.500	Median	:2.0000	Median	:4.087	Median	:5.500
Mean	:4.341	Mean	:2.1856	Mean	:4.186	Mean	:5.527
3rd Qu.	:5.250	3rd Qu.	:2.7143	3rd Qu.	:4.913	3rd Qu.	:6.100
Max.	:7.000	Max.	:6.1429	Max.	:7.435	Max.	:10.000
x9							
Min.	:2.778						
1st Qu.	:4.750						
Median	:5.417						
Mean	:5.374						

```
3rd Qu.:6.083
Max.    :9.250
```

computing the variance-covariance matrix for $P = 9$ variables

```
> N <- nrow(HolzingerSwineford1939)
> S <- cov( HolzingerSwineford1939[, var.names] )
> S <- S * (N-1)/N # ML version
> round(S, 3)
```

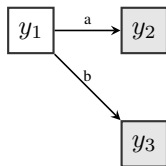
	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	1.358	0.407	0.580	0.505	0.441	0.455	0.085	0.264	0.458
x2	0.407	1.382	0.451	0.209	0.211	0.248	-0.097	0.110	0.244
x3	0.580	0.451	1.275	0.208	0.112	0.244	0.088	0.212	0.374
x4	0.505	0.209	0.208	1.351	1.098	0.896	0.220	0.126	0.243
x5	0.441	0.211	0.112	1.098	1.660	1.015	0.143	0.181	0.295
x6	0.455	0.248	0.244	0.896	1.015	1.196	0.144	0.165	0.236
x7	0.085	-0.097	0.088	0.220	0.143	0.144	1.183	0.535	0.373
x8	0.264	0.110	0.212	0.126	0.181	0.165	0.535	1.022	0.457
x9	0.458	0.244	0.374	0.243	0.295	0.236	0.373	0.457	1.015

the model-implied variance-covariance matrix

- the goal of SEM is to test an a priori specified theory/model, based on empirical data; we would like to know if our model ‘fits’ the data (or not)
- each model can be depicted by a path diagram (we may have several alternative models, each one with its own path diagram)
- each path diagram can be converted to a SEM
- SEM will tell us what the implications are for the data if (assumption!) our model is correct: how ‘should’ the data look like, which patterns should we observe?
- in practice, SEM will tell us how the variance-covariance matrix of the data should look like; we call this the ‘model-implied’ variance-covariance matrix ($\hat{\Sigma}$)
- different models \rightarrow different path diagrams \rightarrow different $\hat{\Sigma}$ matrices
- if $\hat{\Sigma}$ is close to S , the model fits well

example model-implied covariance matrix (1)

- suppose we have three observed (random) variables, y_1 , y_2 and y_3 ; to explain why they are correlated, we may postulate the following model:



$$y_2 = a y_1 + \epsilon_2$$

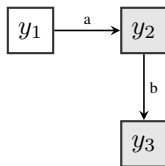
$$y_3 = b y_1 + \epsilon_3$$

- suppose, we set $a = 3$ en $b = 5$, $\text{Var}(y_1) = 10$, $\text{Var}(\epsilon_2) = 20$, $\text{Var}(\epsilon_3) = 30$; then, it can be shown that the model-implied variance-covariance matrix equals

$$\hat{\Sigma} = \begin{bmatrix} 10 & & \\ 30 & 110 & \\ 50 & 150 & 280 \end{bmatrix}$$

example model-implied covariance matrix (2)

- but if we change the path diagram (and keep the parameter values fixed), the model-implied covariance matrix will also change:



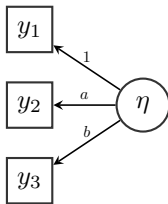
we find

$$\Sigma = \begin{bmatrix} 10 & & \\ 30 & 110 & \\ 150 & 550 & 2780 \end{bmatrix}$$

- two models are said to be *equivalent*, if they imply the same covariance matrix (but note that we did not estimate the parameters here)

example model-implied covariance matrix (3)

- we can also postulate that the correlations among the three observed variables are explained by a common ‘factor’:

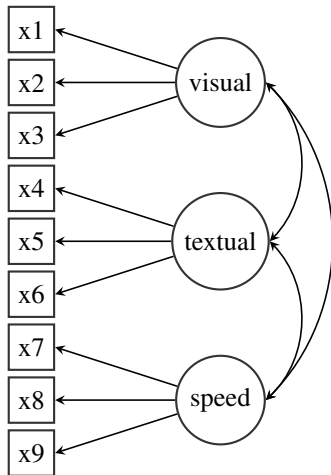


- we find using $\sigma^2(\epsilon_1) = 10$, $\sigma^2(\epsilon_2) = 20$, $\sigma^2(\epsilon_3) = 30$, $\sigma^2(\eta) = 1$:

$$\Sigma = \begin{bmatrix} 11 & & \\ 4 & 36 & \\ 5 & 20 & 55 \end{bmatrix}$$

- we can compare all three $\hat{\Sigma}$ matrices to \mathbf{S} to find out which model fits best

1.3 A first example: a CFA with three factors



- ‘free’ parameters: factor loadings, variances for the factors, covariances between the factors, and residual variances for the indicators

the matrix representation of a CFA model

- the classic LISREL representation uses three model matrices for a CFA
- the LAMBDA matrix contains the ‘factor structure’:

$$\Lambda = \begin{bmatrix} x & 0 & 0 \\ x & 0 & 0 \\ x & 0 & 0 \\ 0 & x & 0 \\ 0 & x & 0 \\ 0 & x & 0 \\ 0 & 0 & x \\ 0 & 0 & x \\ 0 & 0 & x \end{bmatrix}$$

- the variances/covariances of the latent variables are summarized in the PSI matrix:

the standard CFA model: the model implied covariance matrix

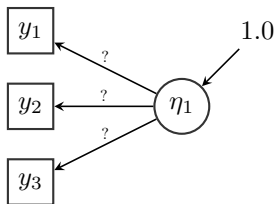
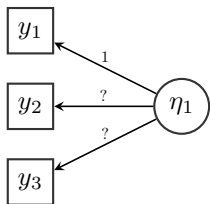
- in the standard CFA model, the ‘implied’ covariance matrix is:

$$\Sigma = \Lambda\Psi\Lambda' + \Theta$$

- all parameters are included in three model matrices
- simple matrix multiplication (and addition) gives us the model implied covariance matrix
- for identification purposes, some parameters need to be fixed to a constant (see next slide)
- estimation problem: choose the ‘free’ parameters, so that the estimated implied covariance matrix ($\hat{\Sigma}$) is ‘as close as possible’ to the observed covariance matrix S
 - generalized (weighted) least-squares estimation (GLS, WLS)
 - maximum likelihood estimation (ML)
 - Bayesian approaches

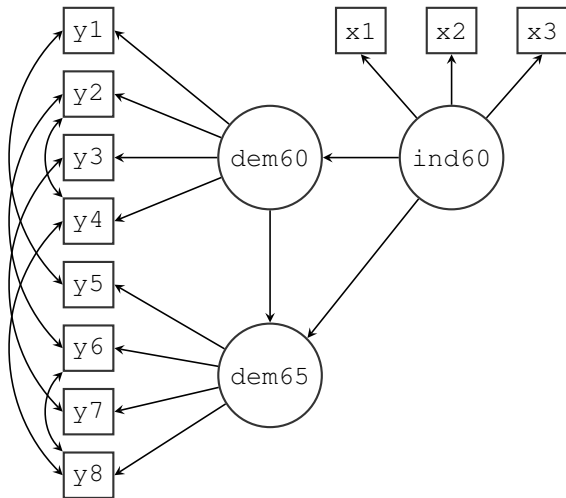
setting the metric of the latent variables: UVI of ULI

1. *Unit Loading Identification (ULI)*:
the factor loading of one (often the first) of the indicators is fixed to 1.0; this indicator is called the *reference* indicator
2. *Unit Variance Identification (UVI)*:
the variance of the factor is fixed to 1.0



- in many models, it does not matter
- in multigroup SEM analysis: we usually use ULI

1.4 A second example: the Political Democracy dataset



model matrices

- this is an example of a ‘full SEM’: the model contains both a measurement part, and a structural part
- we now need 4 model matrices:
 - LAMBDA: the factor loadings
 - THETA: the residual variances (and covariances) of the observed indicators
 - PSI: the (residual) variances and covariances of the latent variables
 - BETA: the regression coefficients of the structural part
- the formula to obtain the model-implied variance-covariance matrix is now slightly more complex:

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

where \mathbf{I} is the identity matrix

1.5 Model estimation

- we seek those values for θ that minimize the difference between what we observe in the data, \mathbf{S} , and what the model implies, $\Sigma(\theta)$
- the final estimated values are denoted by $\hat{\theta}$, and the estimated model-implied covariance matrix can be written as $\hat{\Sigma} = \Sigma(\hat{\theta})$
- there are many ways to quantify this ‘difference’, leading to different discrepancy measures
- the most used discrepancy measure is based on maximum likelihood:

$$F_{ML}(\theta) = \log |\Sigma| + \text{tr}(\mathbf{S}\Sigma^{-1}) - \log |\mathbf{S}| - P$$

- in practice, we replace Σ by $\hat{\Sigma} = \Sigma(\hat{\theta})$
- an alternative is (weighted) least squares, for some weight matrix \mathbf{W} :

$$F_{WLS}(\theta) = (\mathbf{s} - \boldsymbol{\sigma})' \mathbf{W}^{-1} (\mathbf{s} - \boldsymbol{\sigma})$$

where \mathbf{s} and $\boldsymbol{\sigma}$ are the unique elements of \mathbf{S} and Σ respectively

1.6 Model evaluation

evaluation of global fit – chi-square test statistic

- the chi-square test statistic is the primary test of our model
- if the chi-square test statistic is NOT significant, we have a good fit of the model
- this becomes increasingly difficult if the sample size grows

evaluation of global fit – fit indices

- (some) rules of thumb: $CFI/TLI > 0.95$, $RMSEA < 0.05$, $SRMR < 0.06$
- there is a lot of controversy about the use (and misuse) of these fit indices
- a good reference is still Hu & Bentler (1999)
- current practice is to report: chi-square value + df + pvalue, RMSEA, CFI and SRMR (do not cherry pick your fit indices)

1.7 Model respecification

- if the fit of a model is not good, we can adapt (respecify) the model
 - change the number of factors
 - allow for indicators to be related to more than one factor (cross-loadings)
 - allow for correlated residual errors among the observed indicators
 - allow for correlated disturbances among the endogenous latent variables
 - remove problematic indicators . . .
- ideally, all changes should have a sound theoretical justification
- of course, we may let the data speak for itself, and have a look at the modification indices (a more explorative approach)

1.8 Further reading

Kline, R. B. (2015). Principles and practice of structural equation modeling (Fourth Edition). New York: Guilford Press.

Bollen, K.A. (1989). Structural equations with latent variables. New York: Wiley.

Loehlin, J. C., & Beaujean, A. A. (2016). Latent variable models: An introduction to factor, path, and structural equation analysis. Taylor & Francis.

Beaujean, A. A. (2014). Latent variable modeling using R: A step-by-step guide. New York: Routledge.

Finch, W.H., and French, B.F. (2015). Latent Variable Modeling with R. Routledge.

2 Introduction to lavaan

2.1 Software for SEM

software for SEM: commercial – closed-source

- LISREL, EQS, AMOS, MPLUS
- SAS/Stat: proc (T)CALIS, SEPATH (Statistica), RAMONA (Systat), Stata (12 or higher)
- Mx (free, closed-source)

software for SEM: non-commercial – open-source

- outside the R ecosystem: gllamm (Stata), Onyx, semopy, ...
- R packages: sem, OpenMx, lavaan, lava, psychometrics

2.2 The R package ‘lavaan’

what is lavaan?

- **lavaan** is an R package for latent variable analysis:
 - user-friendly syntax, user-friendly interface (the `sem()` function)
 - support for continuous, binary and ordinal data
 - support for missing data, nonnormal data, clustered data, ...
 - available technology: CFA, SEM, path-analysis, growth curve modeling, exploratory factor analysis (EFA), ESEM, multilevel SEM, ...
 - unique for lavaan: bounded estimation, pairwise likelihood (PL) estimation, distributionally weighted least squares (DLS), structural-after-measurement (SAM) approach to SEM, ...
- under development, future plans:
 - technical interface, more algorithms, simulation framework, (multi-group) mixture EFA/CFA/SEM, IRT, new engine, ...

installing lavaan, finding documentation

- **lavaan** depends on the R project for statistical computing:

<https://www.r-project.org>

- to install **lavaan**, simply start up an R session and type:

```
> install.packages("lavaan")
```

- more information about **lavaan**:

<http://lavaan.org>

- the lavaan paper:

Rosseel (2012). lavaan: an R package for structural equation modeling. *Journal of Statistical Software*, 48(2), 1–36.

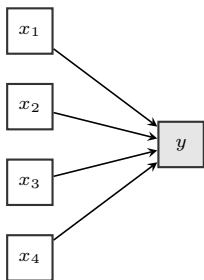
- **lavaan** discussion group (mailing list)

<https://groups.google.com/d/forum/lavaan>

2.3 The lavaan model syntax

using standard R – a simple regression

- using the `lm` function in R:



```
# read in your data
myData <- read.csv("c:/temp/myData.csv")

# fit model using lm
fit <- lm(formula = y ~ x1 + x2 + x3 + x4,
          data = myData)

# show results
summary(fit)
```

- the standard linear model:

$$y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i3} + \beta_4 x_{i4} + \epsilon_i \quad (i = 1, 2, \dots, n)$$

lm() output artificial data (N=100)

```
> summary(fit)
```

```
Call:
```

```
lm(formula = y ~ x1 + x2 + x3 + x4, data = myData)
```

```
Residuals:
```

	Min	1Q	Median	3Q	Max
	-102.372	-29.458	-3.658	27.275	148.404

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	97.7210	4.7200	20.704	<2e-16 ***
x1	5.7733	0.5238	11.022	<2e-16 ***
x2	-1.3214	0.4917	-2.688	0.0085 **
x3	1.1350	0.4575	2.481	0.0149 *
x4	0.2707	0.4779	0.566	0.5724

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

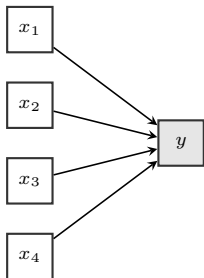
```
Residual standard error: 46.74 on 95 degrees of freedom
```

```
Multiple R-squared:  0.5911,      Adjusted R-squared:  0.5738
```

```
F-statistic: 34.33 on 4 and 95 DF,  p-value: < 2.2e-16
```

the lavaan model syntax – a simple regression

- using lavaan's `sem` function:



```
library(lavaan)
myData <- read.csv("c:/temp/myData.csv")

myModel <- ' y ~ x1 + x2 + x3 + x4 '

# fit model
fit <- sem(model = myModel,
           data = myData)

# show results
summary(fit, nd = 4, header = FALSE)
```

- to 'see' the intercept, use either

```
fit <- sem(model = myModel, data = myData, meanstructure = TRUE)
```

or include it explicitly in the syntax:

```
myModel <- ' y ~ 1 + x1 + x2 + x3 + x4 '
```

(partial) lavaan output**Parameter Estimates:**

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

Regressions:

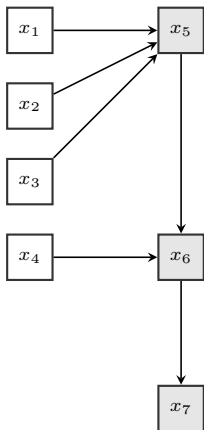
	Estimate	Std.Err	z-value	P(> z)
y ~				
x1	5.7733	0.5105	11.3087	0.0000
x2	-1.3214	0.4792	-2.7574	0.0058
x3	1.1350	0.4459	2.5451	0.0109
x4	0.2707	0.4658	0.5812	0.5611

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y	2075.0999	293.4634	7.0711	0.0000

the lavaan model syntax – path analysis

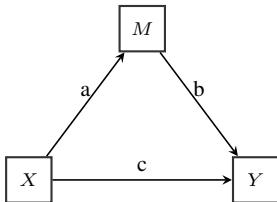
- for each dependent variable, we write a separate regression equation:



```
myModel <- ' x5 ~ x1 + x2 + x3  
            x6 ~ x4 + x5  
            x7 ~ x6  
'
```

the lavaan model syntax – mediation analysis

- a mediation analysis is simple
- we can use labels to refer to specific parameters (here regression coefficients)
- standard errors are based on the bootstrap



```
myModel <- '  
    Y ~ b*M + c*X  
    M ~ a*X  
  
    indirect := a*b  
    total   := c + (a*b)  
,  
  
fit <- sem(model = myModel,  
           data  = myData,  
           se    = "bootstrap")  
  
summary(fit)
```

partial output

Parameter estimates:

Information	Observed
Standard Errors	Bootstrap
Number of requested bootstrap draws	1000
Number of successful bootstrap draws	1000

Regressions:

		Estimate	Std.err	z-value	P(> z)
Y ~					
M	(b)	0.597	0.098	6.068	0.000
X	(c)	2.594	1.210	2.145	0.032
M ~					
X	(a)	2.739	0.999	2.741	0.006

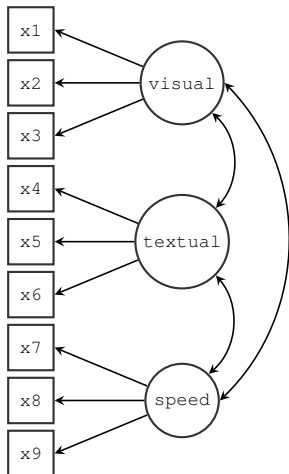
Variances:

	Estimate	Std.err	z-value	P(> z)
.Y	108.700	17.747	6.125	0.000
.M	105.408	16.556	6.367	0.000

Defined parameters:

	Estimate	Std.err	z-value	P(> z)
indirect	1.636	0.645	2.535	0.011
total	4.230	1.383	3.059	0.002

the lavaan model syntax – using `cfa()` or `sem()`

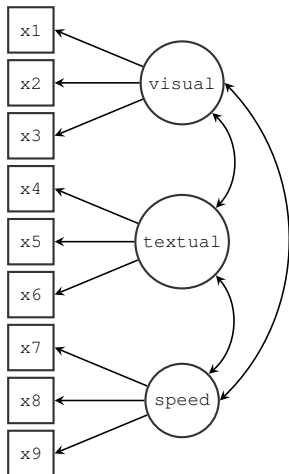


```
HS.model <- ' visual  =~ x1 + x2 + x3
            textual =~ x4 + x5 + x6
            speed   =~ x7 + x8 + x9
            '
```

```
fit <- cfa(model = HS.model,
           data = HolzingerSwineford1939)
```

```
summary(fit, fit.measures = TRUE,
        standardized = TRUE)
```

the lavaan model syntax – using lavaan()



```

HS.model <- '
  # latent variables
  visual  =~ 1*x1 + x2 + x3
  textual =~ 1*x4 + x5 + x6
  speed   =~ 1*x7 + x8 + x9

  # factor (co)variances
  visual  ~~ visual; visual  ~~ textual
  visual  ~~ speed;  textual ~~ textual
  textual ~~ speed;  speed   ~~ speed

  # residual variances
  x1 ~~ x1; x2 ~~ x2; x3 ~~ x3
  x4 ~~ x4; x5 ~~ x5; x6 ~~ x6
  x7 ~~ x7; x8 ~~ x8; x9 ~~ x9
'

fit <- lavaan(model = HS.model,
              data = HolzingerSwineford1939)

summary(fit, fit.measures = TRUE,
        standardized = TRUE)

```

full output

lavaan 0.6-12 ended normally after 35 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	21
Number of observations	301

Model Test User Model:

Test statistic	85.306
Degrees of freedom	24
P-value (Chi-square)	0.000

Model Test Baseline Model:

Test statistic	918.852
Degrees of freedom	36
P-value	0.000

User Model versus Baseline Model:

Comparative Fit Index (CFI)	0.931
Tucker-Lewis Index (TLI)	0.896

Loglikelihood and Information Criteria:

Loglikelihood user model (H0)	-3737.745
Loglikelihood unrestricted model (H1)	-3695.092
Akaike (AIC)	7517.490
Bayesian (BIC)	7595.339
Sample-size adjusted Bayesian (BIC)	7528.739

Root Mean Square Error of Approximation:

RMSEA	0.092
90 Percent confidence interval - lower	0.071
90 Percent confidence interval - upper	0.114
P-value RMSEA \leq 0.05	0.001

Standardized Root Mean Square Residual:

SRMR	0.065
------	-------

Parameter Estimates:

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

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
visual =~						

x1	1.000				0.900	0.772
x2	0.554	0.100	5.554	0.000	0.498	0.424
x3	0.729	0.109	6.685	0.000	0.656	0.581
textual =~						
x4	1.000				0.990	0.852
x5	1.113	0.065	17.014	0.000	1.102	0.855
x6	0.926	0.055	16.703	0.000	0.917	0.838
speed =~						
x7	1.000				0.619	0.570
x8	1.180	0.165	7.152	0.000	0.731	0.723
x9	1.082	0.151	7.155	0.000	0.670	0.665

Covariances:

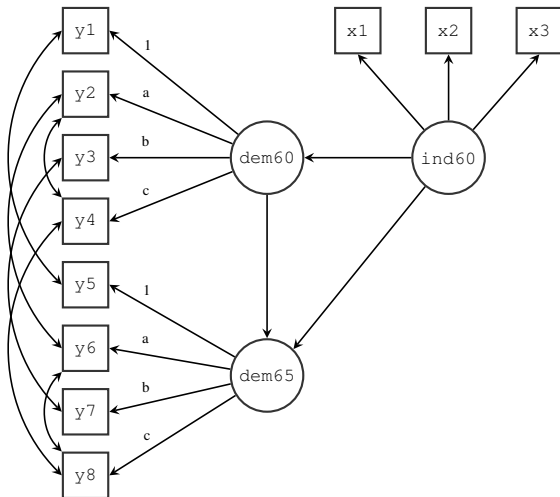
	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
visual ~~						
textual	0.408	0.074	5.552	0.000	0.459	0.459
speed	0.262	0.056	4.660	0.000	0.471	0.471
textual ~~						
speed	0.173	0.049	3.518	0.000	0.283	0.283

Variances:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.x1	0.549	0.114	4.833	0.000	0.549	0.404
.x2	1.134	0.102	11.146	0.000	1.134	0.821
.x3	0.844	0.091	9.317	0.000	0.844	0.662
.x4	0.371	0.048	7.779	0.000	0.371	0.275
.x5	0.446	0.058	7.642	0.000	0.446	0.269
.x6	0.356	0.043	8.277	0.000	0.356	0.298

.x7	0.799	0.081	9.823	0.000	0.799	0.676
.x8	0.488	0.074	6.573	0.000	0.488	0.477
.x9	0.566	0.071	8.003	0.000	0.566	0.558
visual	0.809	0.145	5.564	0.000	1.000	1.000
textual	0.979	0.112	8.737	0.000	1.000	1.000
speed	0.384	0.086	4.451	0.000	1.000	1.000

the lavaan model syntax – equality constraints



fitting the model with lavaan

```
# 1. specifying the model
```

```
model <- '  
  # latent variable definitions  
  ind60 =~ x1 + x2 + x3  
  dem60 =~ y1 + a*y2 + b*y3 + c*y4  
  dem65 =~ y5 + a*y6 + b*y7 + c*y8  
  
  # regressions  
  dem60 ~ ind60  
  dem65 ~ ind60 + dem60  
  
  # residual covariances  
  y1 ~~ y5  
  y2 ~~ y4 + y6  
  y3 ~~ y7  
  y4 ~~ y8  
  y6 ~~ y8  
,
```

```
# 2. fitting the model using the sem() function
```

```
fit <- sem(model, data = PoliticalDemocracy)
```

```
# 3. display the results
```

```
summary(fit, standardized = TRUE)
```

output

lavaan 0.6-12 ended normally after 66 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	31
Number of equality constraints	3
Number of observations	75

Model Test User Model:

Test statistic	40.179
Degrees of freedom	38
P-value (Chi-square)	0.374

Parameter Estimates:

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

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
ind60 =~						
x1	1.000				0.670	0.920
x2	2.180	0.138	15.751	0.000	1.460	0.973

x3		1.818	0.152	11.971	0.000	1.218	0.872
dem60 =~							
y1		1.000				2.201	0.850
y2	(a)	1.191	0.139	8.551	0.000	2.621	0.690
y3	(b)	1.175	0.120	9.755	0.000	2.586	0.758
y4	(c)	1.251	0.117	10.712	0.000	2.754	0.838
dem65 =~							
y5		1.000				2.154	0.817
y6	(a)	1.191	0.139	8.551	0.000	2.565	0.755
y7	(b)	1.175	0.120	9.755	0.000	2.530	0.802
y8	(c)	1.251	0.117	10.712	0.000	2.694	0.829

Regressions:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
dem60 ~						
ind60	1.471	0.392	3.750	0.000	0.448	0.448
dem65 ~						
ind60	0.600	0.226	2.661	0.008	0.187	0.187
dem60	0.865	0.075	11.554	0.000	0.884	0.884

Covariances:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.y1 ~~						
.y5	0.583	0.356	1.637	0.102	0.583	0.281
.y2 ~~						
.y4	1.440	0.689	2.092	0.036	1.440	0.291
.y6	2.183	0.737	2.960	0.003	2.183	0.356
.y3 ~~						

.y7	0.712	0.611	1.165	0.244	0.712	0.169
.y4 ~~						
.y8	0.363	0.444	0.817	0.414	0.363	0.111
.y6 ~~						
.y8	1.372	0.577	2.378	0.017	1.372	0.338

Variances:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.x1	0.081	0.019	4.182	0.000	0.081	0.154
.x2	0.120	0.070	1.729	0.084	0.120	0.053
.x3	0.467	0.090	5.177	0.000	0.467	0.239
.y1	1.855	0.433	4.279	0.000	1.855	0.277
.y2	7.581	1.366	5.549	0.000	7.581	0.525
.y3	4.956	0.956	5.182	0.000	4.956	0.426
.y4	3.225	0.723	4.458	0.000	3.225	0.298
.y5	2.313	0.479	4.831	0.000	2.313	0.333
.y6	4.968	0.921	5.393	0.000	4.968	0.430
.y7	3.560	0.710	5.018	0.000	3.560	0.357
.y8	3.308	0.704	4.701	0.000	3.308	0.313
ind60	0.449	0.087	5.175	0.000	1.000	1.000
.dem60	3.875	0.866	4.477	0.000	0.800	0.800
.dem65	0.164	0.227	0.725	0.469	0.035	0.035

2.4 lavaan: a brief user's guide

example: fitted()

```
> fit <- cfa(HS.model, data = HolzingerSwineford1939)
> fitted(fit)
```

```
$cov
  x1    x2    x3    x4    x5    x6    x7    x8    x9
x1 1.358
x2 0.448 1.382
x3 0.590 0.327 1.275
x4 0.408 0.226 0.298 1.351
x5 0.454 0.252 0.331 1.090 1.660
x6 0.378 0.209 0.276 0.907 1.010 1.196
x7 0.262 0.145 0.191 0.173 0.193 0.161 1.183
x8 0.309 0.171 0.226 0.205 0.228 0.190 0.453 1.022
x9 0.284 0.157 0.207 0.188 0.209 0.174 0.415 0.490 1.015
```

example: lavInspect()

```
> lavInspect (fit)
```

```
$lambda
```

	visual	textul	speed
x1	0	0	0
x2	1	0	0
x3	2	0	0
x4	0	0	0
x5	0	3	0
x6	0	4	0
x7	0	0	0
x8	0	0	5
x9	0	0	6

```
$theta
```

	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	7								
x2	0	8							
x3	0	0	9						
x4	0	0	0	10					
x5	0	0	0	0	11				
x6	0	0	0	0	0	12			
x7	0	0	0	0	0	0	13		
x8	0	0	0	0	0	0	0	14	
x9	0	0	0	0	0	0	0	0	15


```
$psi
      visual textual speed
visual  16
textual 19      17
speed   20      21      18
```

```
> lavInspect(fit, "sampstat")
```

```
$cov
      x1      x2      x3      x4      x5      x6      x7      x8      x9
x1  1.358
x2  0.407  1.382
x3  0.580  0.451  1.275
x4  0.505  0.209  0.208  1.351
x5  0.441  0.211  0.112  1.098  1.660
x6  0.455  0.248  0.244  0.896  1.015  1.196
x7  0.085 -0.097  0.088  0.220  0.143  0.144  1.183
x8  0.264  0.110  0.212  0.126  0.181  0.165  0.535  1.022
x9  0.458  0.244  0.374  0.243  0.295  0.236  0.373  0.457  1.015
```

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

```
      visual textual speed
visual  0.809
textual 0.408  0.979
speed   0.262  0.173  0.384
```

```
> lavTech(fit, "cov.lv")
```

```
[[1]]
      [,1]      [,2]      [,3]
[1,] 0.8093160 0.4082324 0.2622246
[2,] 0.4082324 0.9794914 0.1734947
[3,] 0.2622246 0.1734947 0.3837476
```

```
> lavTech(fit, "cov.lv", add.labels = TRUE, drop.list.single.group = TRUE)
```

```
      visual  textual  speed
visual 0.8093160 0.4082324 0.2622246
textual 0.4082324 0.9794914 0.1734947
speed   0.2622246 0.1734947 0.3837476
```

example: fitMeasures()

```
> fitMeasures(fit)
```

```

      npar              fmin              chisq              df
21.000              0.142             85.306             24.000
pvalue      baseline.chisq      baseline.df      baseline.pvalue
0.000              918.852             36.000             0.000
      cfi              tli              nnfi              rfi
0.931              0.896             0.896             0.861
      nfi              pnfi              ifi              rni
0.907              0.605             0.931             0.931
      logl      unrestricted.logl              aic              bic
-3737.745      -3695.092             7517.490             7595.339
      ntotal              bic2              rmsea      rmsea.ci.lower
301.000              7528.739             0.092             0.071
rmsea.ci.upper      rmsea.pvalue              rmr              rmr_nomean
0.114              0.001             0.082             0.082
      srmr      srmr_bentler      srmr_bentler_nomean      crmr
0.065              0.065             0.065             0.073
cmrm_nomean      srmr_mplus      srmr_mplus_nomean      cn_05
0.073              0.065             0.065             129.490
      cn_01              gfi              agfi              pgfi
152.654              0.943             0.894             0.503
      mfi              ecvi
0.903              0.423

```

example: parameterTable()

```
> parameterTable(fit)[1:21, 1:13]
```

	id	lhs	op	rhs	user	block	group	free	ustart	exo	label	plabel	start
1	1	visual	=~	x1	1	1	1	0	1	0		.p1.	1.000
2	2	visual	=~	x2	1	1	1	1	NA	0		.p2.	0.778
3	3	visual	=~	x3	1	1	1	2	NA	0		.p3.	1.107
4	4	textual	=~	x4	1	1	1	0	1	0		.p4.	1.000
5	5	textual	=~	x5	1	1	1	3	NA	0		.p5.	1.133
6	6	textual	=~	x6	1	1	1	4	NA	0		.p6.	0.924
7	7	speed	=~	x7	1	1	1	0	1	0		.p7.	1.000
8	8	speed	=~	x8	1	1	1	5	NA	0		.p8.	1.225
9	9	speed	=~	x9	1	1	1	6	NA	0		.p9.	0.854
10	10	x1	~~	x1	0	1	1	7	NA	0		.p10.	0.679
11	11	x2	~~	x2	0	1	1	8	NA	0		.p11.	0.691
12	12	x3	~~	x3	0	1	1	9	NA	0		.p12.	0.637
13	13	x4	~~	x4	0	1	1	10	NA	0		.p13.	0.675
14	14	x5	~~	x5	0	1	1	11	NA	0		.p14.	0.830
15	15	x6	~~	x6	0	1	1	12	NA	0		.p15.	0.598
16	16	x7	~~	x7	0	1	1	13	NA	0		.p16.	0.592
17	17	x8	~~	x8	0	1	1	14	NA	0		.p17.	0.511
18	18	x9	~~	x9	0	1	1	15	NA	0		.p18.	0.508
19	19	visual	~~	visual	0	1	1	16	NA	0		.p19.	0.050
20	20	textual	~~	textual	0	1	1	17	NA	0		.p20.	0.050
21	21	speed	~~	speed	0	1	1	18	NA	0		.p21.	0.050

example: parameterEstimates()

```
> parameterEstimates(fit)[1:21,]
```

	lhs	op	rhs	est	se	z	pvalue	ci.lower	ci.upper
1	visual	=~	x1	1.000	0.000	NA	NA	1.000	1.000
2	visual	=~	x2	0.554	0.100	5.554	0	0.358	0.749
3	visual	=~	x3	0.729	0.109	6.685	0	0.516	0.943
4	textual	=~	x4	1.000	0.000	NA	NA	1.000	1.000
5	textual	=~	x5	1.113	0.065	17.014	0	0.985	1.241
6	textual	=~	x6	0.926	0.055	16.703	0	0.817	1.035
7	speed	=~	x7	1.000	0.000	NA	NA	1.000	1.000
8	speed	=~	x8	1.180	0.165	7.152	0	0.857	1.503
9	speed	=~	x9	1.082	0.151	7.155	0	0.785	1.378
10	x1	~~	x1	0.549	0.114	4.833	0	0.326	0.772
11	x2	~~	x2	1.134	0.102	11.146	0	0.934	1.333
12	x3	~~	x3	0.844	0.091	9.317	0	0.667	1.022
13	x4	~~	x4	0.371	0.048	7.779	0	0.278	0.465
14	x5	~~	x5	0.446	0.058	7.642	0	0.332	0.561
15	x6	~~	x6	0.356	0.043	8.277	0	0.272	0.441
16	x7	~~	x7	0.799	0.081	9.823	0	0.640	0.959
17	x8	~~	x8	0.488	0.074	6.573	0	0.342	0.633
18	x9	~~	x9	0.566	0.071	8.003	0	0.427	0.705
19	visual	~~	visual	0.809	0.145	5.564	0	0.524	1.094
20	textual	~~	textual	0.979	0.112	8.737	0	0.760	1.199
21	speed	~~	speed	0.384	0.086	4.451	0	0.215	0.553

example: modindices()

```
> modindices(fit, sort = TRUE, minimum.value = 5)
```

	lhs	op	rhs	mi	epc	sepc.lv	sepc.all	sepc.nox
30	visual	=~	x9	36.411	0.577	0.519	0.515	0.515
76	x7	~~	x8	34.145	0.536	0.536	0.859	0.859
28	visual	=~	x7	18.631	-0.422	-0.380	-0.349	-0.349
78	x8	~~	x9	14.946	-0.423	-0.423	-0.805	-0.805
33	textual	=~	x3	9.151	-0.272	-0.269	-0.238	-0.238
55	x2	~~	x7	8.918	-0.183	-0.183	-0.192	-0.192
31	textual	=~	x1	8.903	0.350	0.347	0.297	0.297
51	x2	~~	x3	8.532	0.218	0.218	0.223	0.223
59	x3	~~	x5	7.858	-0.130	-0.130	-0.212	-0.212
26	visual	=~	x5	7.441	-0.210	-0.189	-0.147	-0.147
50	x1	~~	x9	7.335	0.138	0.138	0.247	0.247
65	x4	~~	x6	6.220	-0.235	-0.235	-0.646	-0.646
66	x4	~~	x7	5.920	0.098	0.098	0.180	0.180
48	x1	~~	x7	5.420	-0.129	-0.129	-0.195	-0.195
77	x7	~~	x9	5.183	-0.187	-0.187	-0.278	-0.278

example: lavResiduals()

```
> lavResiduals(fit)
```

```
$type
```

```
[1] "cor.bentler"
```

```
$cov
```

	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	0.000								
x2	-0.030	0.000							
x3	-0.008	0.094	0.000						
x4	0.071	-0.012	-0.068	0.000					
x5	-0.009	-0.027	-0.151	0.005	0.000				
x6	0.060	0.030	-0.026	-0.009	0.003	0.000			
x7	-0.140	-0.189	-0.084	0.037	-0.036	-0.014	0.000		
x8	-0.039	-0.052	-0.012	-0.067	-0.036	-0.022	0.075	0.000	
x9	0.149	0.073	0.147	0.048	0.067	0.056	-0.038	-0.032	0.000

```
$cov.z
```

	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	0.000								
x2	-1.996	0.000							
x3	-0.997	2.689	0.000						
x4	2.679	-0.284	-1.899	0.000					
x5	-0.359	-0.591	-4.157	1.545	0.000				
x6	2.155	0.681	-0.711	-2.588	0.942	0.000			
x7	-3.773	-3.654	-1.858	0.865	-0.842	-0.326	0.000		

```
x8 -1.380 -1.119 -0.300 -2.021 -1.099 -0.641 4.823 0.000
x9 4.077 1.606 3.518 1.225 1.701 1.423 -2.325 -4.132 0.000
```

```
$summary
```

```
                cov
srmr            0.065
srmr.se         0.006
srmr.exactfit.z 6.063
srmr.exactfit.pvalue 0.000
usrmr           0.058
usrmr.se        0.010
usrmr.ci.lower  0.042
usrmr.ci.upper  0.074
usrmr.closefit.h0.value 0.050
usrmr.closefit.z 0.832
usrmr.closefit.pvalue 0.203
```


example: lavTestLRT()

```
> fit0 <- update(fit, orthogonal = TRUE)
> lavTestLRT(fit0, fit)
```

Chi-Squared Difference Test

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit	24	7517.5	7595.3	85.305			
fit0	27	7579.7	7646.4	153.527	68.222	3	1.026e-14 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

2.5 Meanstructures

adding the means in lavaan

- when the `meanstructure` argument is set to `TRUE`, a meanstructure is added to the model

```
> fit <- cfa(HS.model, data = HolzingerSwineford1939,  
+           meanstructure = TRUE)
```

- if no restrictions are imposed on the means, the fit will be identical to the non-meanstructure fit
- we add p datapoints (the mean vector)
- we add p free parameters (the intercepts of the observed variables)
- we fix the latent means to zero
- the number of degrees of freedom does not change

output meanstructure = TRUE

lavaan 0.6-12 ended normally after 35 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	30
Number of observations	301

Model Test User Model:

Test statistic	85.306
Degrees of freedom	24
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)
visual =~				
x1	1.000			
x2	0.554	0.100	5.554	0.000
x3	0.729	0.109	6.685	0.000

```

textual =~
  x4      1.000
  x5      1.113    0.065    17.014    0.000
  x6      0.926    0.055    16.703    0.000
speed =~
  x7      1.000
  x8      1.180    0.165     7.152    0.000
  x9      1.082    0.151     7.155    0.000

```

Covariances:

	Estimate	Std.Err	z-value	P(> z)
visual ~~				
textual	0.408	0.074	5.552	0.000
speed	0.262	0.056	4.660	0.000
textual ~~				
speed	0.173	0.049	3.518	0.000

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
.x1	4.936	0.067	73.473	0.000
.x2	6.088	0.068	89.855	0.000
.x3	2.250	0.065	34.579	0.000
.x4	3.061	0.067	45.694	0.000
.x5	4.341	0.074	58.452	0.000
.x6	2.186	0.063	34.667	0.000
.x7	4.186	0.063	66.766	0.000
.x8	5.527	0.058	94.854	0.000
.x9	5.374	0.058	92.546	0.000

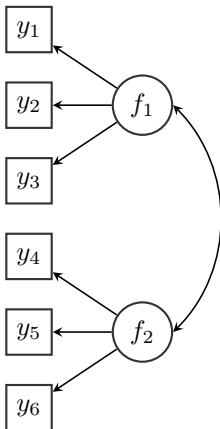
visual	0.000
textual	0.000
speed	0.000

Variances:

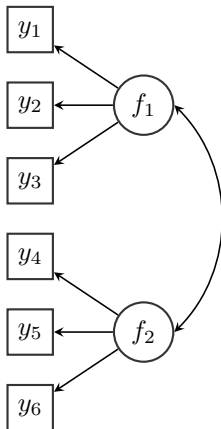
	Estimate	Std.Err	z-value	P(> z)
.x1	0.549	0.114	4.833	0.000
.x2	1.134	0.102	11.146	0.000
.x3	0.844	0.091	9.317	0.000
.x4	0.371	0.048	7.779	0.000
.x5	0.446	0.058	7.642	0.000
.x6	0.356	0.043	8.277	0.000
.x7	0.799	0.081	9.823	0.000
.x8	0.488	0.074	6.573	0.000
.x9	0.566	0.071	8.003	0.000
visual	0.809	0.145	5.564	0.000
textual	0.979	0.112	8.737	0.000
speed	0.384	0.086	4.451	0.000

2.6 Multiple groups

GROUP 1



GROUP 2

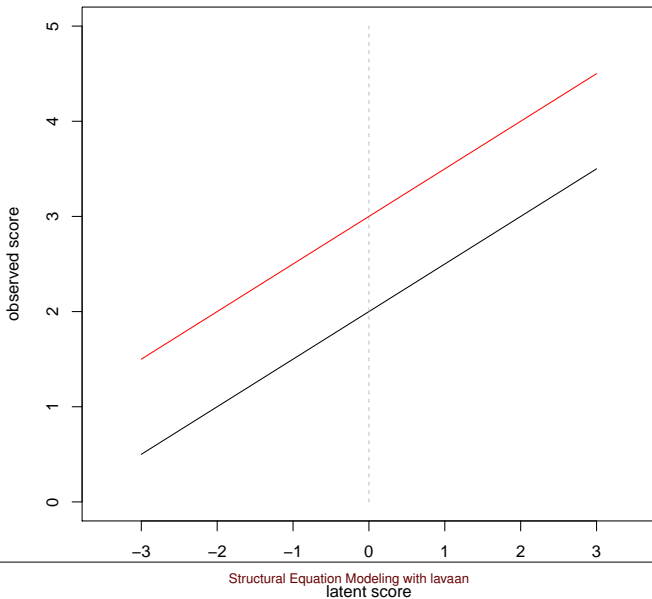


- can we compare the means of the latent variables?

2.7 Measurement invariance

- we can only compare the means of the latent variables across groups if ‘measurement invariance’ across groups has been established
- testing for measurement invariance involves a fixed sequence of model comparison tests
- one typical sequence involves 3 steps:
 1. Model 1: configural invariance. The same factor structure is imposed on all groups.
 2. Model 2: weak invariance. The factor loadings are constrained to be equal across groups.
 3. Model 3: strong invariance. The factor loadings and intercepts are constrained to be equal across groups.
- other sequences involve more steps; for example ‘strict invariance’ implies constraining the residual variances too

example weak invariance (two groups)



measurement invariance in lavaan - using the group.equal argument

- step 1: fit the configural invariance model (fit1)

```
> fit1 <- cfa(HS.model, data = HolzingerSwineford1939, group = "school")  
> fitMeasures(fit1, c("chisq", "df", "pvalue", "cfi", "rmsea", "srmr"))
```

chisq	df	pvalue	cfi	rmsea	srmr
115.851	48.000	0.000	0.923	0.097	0.068

- step 2: fit the weak invariance model (fit2)

```
> fit2 <- cfa(HS.model, data = HolzingerSwineford1939, group = "school",  
+           group.equal = "loadings")  
> fitMeasures(fit2, c("chisq", "df", "pvalue", "cfi", "rmsea", "srmr"))
```

chisq	df	pvalue	cfi	rmsea	srmr
124.044	54.000	0.000	0.921	0.093	0.072

- step 2b: compare with configural invariance model

```
> anova(fit1, fit2)
```

Chi-Squared Difference Test

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit1	48	7484.4	7706.8	115.85			
fit2	54	7480.6	7680.8	124.04	8.1922	6	0.2244

- step 3: fit the strong invariance model (fit3)

```
> fit3 <- cfa(HS.model, data = HolzingerSwineford1939, group = "school",
+           group.equal = c("loadings", "intercepts"))
> fitMeasures(fit3, c("chisq", "df", "pvalue", "cfi", "rmsea", "srmr"))
```

chisq	df	pvalue	cfi	rmsea	srmr
164.103	60.000	0.000	0.882	0.107	0.082

- step 3a: compare with weak invariance model

```
> anova(fit2, fit3)
```

Chi-Squared Difference Test

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit2	54	7480.6	7680.8	124.04			
fit3	60	7508.6	7686.6	164.10	40.059	6	4.435e-07 ***

 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

output strong invariance model

lavaan 0.6-12 ended normally after 60 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	63
Number of equality constraints	15
Number of observations per group:	
Pasteur	156
Grant-White	145

Model Test User Model:

Test statistic	164.103
Degrees of freedom	60
P-value (Chi-square)	0.000
Test statistic for each group:	
Pasteur	90.210
Grant-White	73.892

Parameter Estimates:

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

Group 1 [Pasteur]:

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
visual =~				
x1	1.000			
x2 (.p2.)	0.576	0.101	5.713	0.000
x3 (.p3.)	0.798	0.112	7.146	0.000
textual =~				
x4	1.000			
x5 (.p5.)	1.120	0.066	16.965	0.000
x6 (.p6.)	0.932	0.056	16.608	0.000
speed =~				
x7	1.000			
x8 (.p8.)	1.130	0.145	7.786	0.000
x9 (.p9.)	1.009	0.132	7.667	0.000

Covariances:

	Estimate	Std.Err	z-value	P(> z)
visual ~~				
textual	0.410	0.095	4.293	0.000
speed	0.178	0.066	2.687	0.007
textual ~~				
speed	0.180	0.062	2.900	0.004

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
--	----------	---------	---------	---------

.x1	(.25.)	5.001	0.090	55.760	0.000
.x2	(.26.)	6.151	0.077	79.905	0.000
.x3	(.27.)	2.271	0.083	27.387	0.000
.x4	(.28.)	2.778	0.087	31.953	0.000
.x5	(.29.)	4.035	0.096	41.858	0.000
.x6	(.30.)	1.926	0.079	24.426	0.000
.x7	(.31.)	4.242	0.073	57.975	0.000
.x8	(.32.)	5.630	0.072	78.531	0.000
.x9	(.33.)	5.465	0.069	79.016	0.000
visual		0.000			
textual		0.000			
speed		0.000			

Variances:

	Estimate	Std.Err	z-value	P(> z)
.x1	0.555	0.139	3.983	0.000
.x2	1.296	0.158	8.186	0.000
.x3	0.944	0.136	6.929	0.000
.x4	0.445	0.069	6.430	0.000
.x5	0.502	0.082	6.136	0.000
.x6	0.263	0.050	5.264	0.000
.x7	0.888	0.120	7.416	0.000
.x8	0.541	0.095	5.706	0.000
.x9	0.654	0.096	6.805	0.000
visual	0.796	0.172	4.641	0.000
textual	0.879	0.131	6.694	0.000
speed	0.322	0.082	3.914	0.000

Group 2 [Grant-White]:

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
visual =~				
x1	1.000			
x2 (.p2.)	0.576	0.101	5.713	0.000
x3 (.p3.)	0.798	0.112	7.146	0.000
textual =~				
x4	1.000			
x5 (.p5.)	1.120	0.066	16.965	0.000
x6 (.p6.)	0.932	0.056	16.608	0.000
speed =~				
x7	1.000			
x8 (.p8.)	1.130	0.145	7.786	0.000
x9 (.p9.)	1.009	0.132	7.667	0.000

Covariances:

	Estimate	Std.Err	z-value	P(> z)
visual ~~				
textual	0.427	0.097	4.417	0.000
speed	0.329	0.082	4.006	0.000
textual ~~				
speed	0.236	0.073	3.224	0.001

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
--	----------	---------	---------	---------

.x1	(.25.)	5.001	0.090	55.760	0.000
.x2	(.26.)	6.151	0.077	79.905	0.000
.x3	(.27.)	2.271	0.083	27.387	0.000
.x4	(.28.)	2.778	0.087	31.953	0.000
.x5	(.29.)	4.035	0.096	41.858	0.000
.x6	(.30.)	1.926	0.079	24.426	0.000
.x7	(.31.)	4.242	0.073	57.975	0.000
.x8	(.32.)	5.630	0.072	78.531	0.000
.x9	(.33.)	5.465	0.069	79.016	0.000
visual		-0.148	0.122	-1.211	0.226
textual		0.576	0.117	4.918	0.000
speed		-0.177	0.090	-1.968	0.049

Variances:

	Estimate	Std.Err	z-value	P(> z)
.x1	0.654	0.128	5.094	0.000
.x2	0.964	0.123	7.812	0.000
.x3	0.641	0.101	6.316	0.000
.x4	0.343	0.062	5.534	0.000
.x5	0.376	0.073	5.133	0.000
.x6	0.437	0.067	6.559	0.000
.x7	0.625	0.095	6.574	0.000
.x8	0.434	0.088	4.914	0.000
.x9	0.522	0.086	6.102	0.000
visual	0.708	0.160	4.417	0.000
textual	0.870	0.131	6.659	0.000
speed	0.505	0.115	4.379	0.000

2.8 Missing data

missing data in lavaan

- in lavaan 0.6, the default is listwise deletion (but this may change in future versions)
 - the goal is to alert the user that data is missing
 - the user should examine: which variables have many missing values?
is there a pattern?
- available approaches in lavaan:
 - ‘full information’ ML (`missing = "fiml"`)
 - two-stage approach (`missing = "two.stage"`)
- multiple imputation in lavaan:
 - create imputed datasets (eg., using the `mice` package) + `lavaanList()`
 - the `runMI()` function in the `semTools` package

example: lavaan + listwise

```
> fit <- cfa(HS.model, data = HS.missing, missing = "listwise")
> fit
```

lavaan 0.6-12 ended normally after 36 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	21	
	Used	Total
Number of observations	259	301

Model Test User Model:

Test statistic	89.144
Degrees of freedom	24
P-value (Chi-square)	0.000

example: lavaan + fiml

```
> fit <- cfa(HS.model, data = HS.missing, missing = "fiml") # or missing = "ml"  
> fit
```

lavaan 0.6-12 ended normally after 54 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	30
Number of observations	301
Number of missing patterns	13

Model Test User Model:

Test statistic	86.624
Degrees of freedom	24
P-value (Chi-square)	0.000

example: lavaan + two.stage

```
> fit <- cfa(HS.model, data = HS.missing, missing = "two.stage")
> fit
```

lavaan 0.6-12 ended normally after 37 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	30
Number of observations	301
Number of missing patterns	13

Model Test User Model:

	Standard	Robust
Test Statistic	91.404	87.698
Degrees of freedom	24	24
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.042
Satorra-Bentler correction		

- a robust test statistic and robust standard errors are needed to take the two-stage estimation process into account
- outperforms ‘fiml’ in the non-normal case (see Savalei & Falk, 2014)

2.9 Nonnormal data and alternative estimators

what if the data are NOT normally distributed?

- in the real world, data may never be normally distributed
- two types:
 - categorical and/or limited-dependent outcomes: binary, ordinal, nominal, counts, censored (WLSMV, logit/probit)
 - continuous outcomes, not normally distributed: skewed, too flat/too peaked (kurtosis), ...
- three strategies to deal with continuous non-normal data
 1. asymptotically distribution-free estimation
 2. ML estimation with ‘robust’ standard errors, and a ‘robust’ test statistic for model evaluation
 3. bootstrapping

robust method 1: asymptotically distribution-free (ADF) estimation

- the ADF estimator (Browne, 1984) makes no assumption of normality and is part of a larger family of estimators called weighted least squares (WLS) estimators:

$$F_{WLS} = (\mathbf{s} - \hat{\boldsymbol{\sigma}})^\top \mathbf{W}^{-1} (\mathbf{s} - \hat{\boldsymbol{\sigma}})$$

where \mathbf{s} and $\hat{\boldsymbol{\sigma}}$ are vectors containing the non-duplicated elements in the sample (\mathbf{S}) and model-implied ($\hat{\boldsymbol{\Sigma}}$) covariance matrix respectively

- the weight matrix \mathbf{W} utilized with the ADF estimator is the asymptotic covariance matrix: a matrix of the covariances of the observed sample variances and covariances
- unfortunately, empirical research has shown that the ADF method breaks down unless the sample size is huge (e.g., $N > 5000$)
- in lavaan:

```
fit <- cfa(HS.model, data = HolzingerSwineford1939, estimator = "WLS")
```

robust method 2: robust ML

1. parameter estimates: vanilla ML

- if ML is used, the parameter estimates are still consistent (if the model is identified and correctly specified)

2. 'robust' standard errors

- if data is non-normal, the standard errors tend to be too small (as much as 25-50%)
- 'robust' standard errors correct for non-normality

3. 'robust' scaled (chi-square) test statistic

- if data is non-normal, the usual model (chi-square) test statistic tends to be too large
- the **Satorra-Bentler scaled test statistic** rescales the value of the ML-based chi-square test statistic by an amount that reflects the degree of kurtosis

robust ML in lavaan

- robust standard errors

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          se = "robust")
```

- Satorra-Bentler scaled test statistic

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          test = "Satorra-Bentler")
```

- robust standard errors + scaled test statistic

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          se = "robust", test = "Satorra-Bentler")
```

- estimator MLM = robust standard errors + scaled test statistic

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          estimator = "MLM")
```

- alternative: estimator MLR (also for missing data)

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          estimator = "MLR", missing = "ml")
```

output: robust standard errors and scaled test statistic (MLM)

```
> fit <- cfa(HS.model, data = HolzingerSwineford1939, estimator = "MLM")
> summary(fit)
```

lavaan 0.6-12 ended normally after 35 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	21
Number of observations	301

Model Test User Model:

	Standard	Robust
Test Statistic	85.306	80.872
Degrees of freedom	24	24
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.055
Satorra-Bentler correction		

Parameter Estimates:

Standard errors	Robust.sem
Information	Expected
Information saturated (h1) model	Structured

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
visual =~				
x1	1.000			
x2	0.554	0.103	5.359	0.000
x3	0.729	0.115	6.367	0.000
textual =~				
x4	1.000			
x5	1.113	0.066	16.762	0.000
x6	0.926	0.060	15.497	0.000
speed =~				
x7	1.000			
x8	1.180	0.152	7.758	0.000
x9	1.082	0.132	8.169	0.000

Covariances:

	Estimate	Std.Err	z-value	P(> z)
visual ~~				
textual	0.408	0.082	4.966	0.000
speed	0.262	0.055	4.762	0.000
textual ~~				
speed	0.173	0.055	3.139	0.002

Variances:

	Estimate	Std.Err	z-value	P(> z)
.x1	0.549	0.138	3.968	0.000
.x2	1.134	0.107	10.554	0.000
.x3	0.844	0.085	9.985	0.000
.x4	0.371	0.050	7.423	0.000

.x5	0.446	0.058	7.688	0.000
.x6	0.356	0.046	7.700	0.000
.x7	0.799	0.079	10.168	0.000
.x8	0.488	0.074	6.567	0.000
.x9	0.566	0.068	8.332	0.000
visual	0.809	0.167	4.837	0.000
textual	0.979	0.121	8.109	0.000
speed	0.384	0.083	4.635	0.000

robust method 3: bootstrapping

- bootstrapping standard errors:

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          se = "bootstrap", verbose = TRUE, bootstrap = 1000)
```

- bootstrapping the test statistic

```
fit <- cfa(HS.model, data = HolzingerSwineford1939,  
          test = "bootstrap", verbose = TRUE, bootstrap = 1000)
```

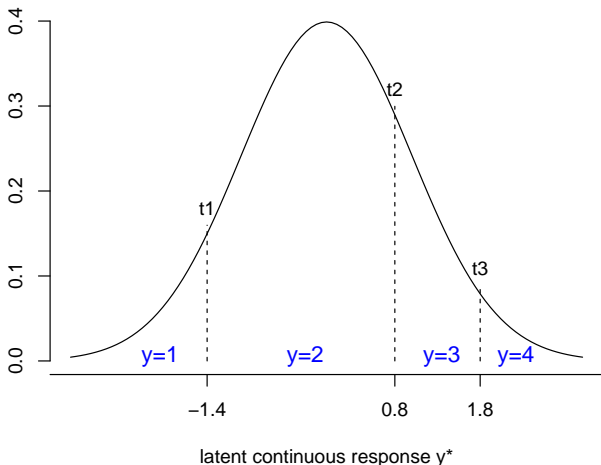
- when we use `se = "bootstrap"`, the `parameterEstimates()` output will contain bootstrap based confidence intervals

2.10 Categorical data

- limited information approach
 - only univariate and bivariate information is used
 - estimation often proceeds in two or three stages; the first stages use maximum likelihood, the last stage uses (weighted) least squares
 - mainly developed in the SEM literature
 - perhaps the best known implementation is in Mplus (WLSMV)
- full information approach
 - all information is used
 - most practical: marginal maximum likelihood estimation
 - requires numerical integration (number of dimensions = number of latent variables)
 - mainly developed in the IRT literature (and GLMM literature)
 - only recently incorporated in modern SEM software

WLS(MV) approach: stage 1 – estimating the thresholds

- an observed variable y can often be viewed as a partial observation of a latent continuous response y^* ; eg ordinal variable with $K = 4$ response categories:



stage 2 – estimating tetrachoric, polychoric, . . . , correlations

- estimate tetrachoric/polychoric/. . . correlation from bivariate data:
 - tetrachoric (binary – binary)
 - polychoric (ordered – ordered)
 - polyserial (ordered – numeric)
 - biserial (binary – numeric)
 - pearson (numeric – numeric)
- ML estimation is available (see eg. Olsson 1979 and 1982)
 - two-step: first estimate thresholds using univariate information only; then, keeping the thresholds fixed, estimate the correlation
 - one-step: estimate thresholds and correlation simultaneously
- if exogenous covariates are involved, the correlations are based on the residual values of y^* (eg bivariate probit regression)

stage 3 – estimating the SEM model

- third stage uses weighted least squares:

$$F_{WLS} = (\mathbf{s} - \hat{\boldsymbol{\sigma}})^\top \mathbf{W}^{-1} (\mathbf{s} - \hat{\boldsymbol{\sigma}})$$

where \mathbf{s} and $\hat{\boldsymbol{\sigma}}$ are vectors containing all relevant sample-based and model-based statistics respectively

- \mathbf{s} contains: thresholds, correlations, optionally regression slopes of exogenous covariates, optionally variances and means of continuous variables
- the weight matrix \mathbf{W} is (a consistent estimator of) the asymptotic covariance matrix of the sample statistics (\mathbf{s})
- robust version: WLSMV
 - use the diagonal of \mathbf{W} only for estimation (DWLS)
 - use the full matrix for inference (standard errors and test statistic)
 - ‘MV’ stands for the Satterthwaite’s mean and variance corrected test statistic

example

```
> # binary version of Holzinger & Swineford
> HS9 <- HolzingerSwineford1939[,c("x1", "x2", "x3", "x4", "x5",
+                                "x6", "x7", "x8", "x9")]
> HSbinary <- as.data.frame( lapply(HS9, cut, 2, labels = FALSE) )

> # single factor model
> model <- ' visual  =~ x1 + x2 + x3
+          textual  =~ x4 + x5 + x6
+          speed    =~ x7 + x8 + x9 '

> # method 1: list all the `ordered` observed variables
> fit <- cfa(model, data = HSbinary, ordered = c("x1", "x2", "x3", "x4", "x5",
+                                               "x6", "x7", "x8", "x9"))
> #
> # method 2: these are also the names of the data.frame `HSbinary`
> fit <- cfa(model, data = HSbinary, ordered = names(HSbinary))
> #
> # method 3: because ALL observed variables are ordered, we can use the shortcut:
> fit <- cfa(model, data = HSbinary, ordered = TRUE)
```


output

```
> summary(fit, fit.measures = TRUE, standardized = TRUE)
```

```
lavaan 0.6-12 ended normally after 35 iterations
```

Estimator	DWLS
Optimization method	NLMINB
Number of model parameters	21
Number of observations	301

```
Model Test User Model:
```

	Standard	Robust
Test Statistic	30.918	38.427
Degrees of freedom	24	24
P-value (Chi-square)	0.156	0.031
Scaling correction factor		0.869
Shift parameter		2.861
simple second-order correction		

```
Model Test Baseline Model:
```

Test statistic	582.533	468.233
Degrees of freedom	36	36
P-value	0.000	0.000
Scaling correction factor		1.264

User Model versus Baseline Model:

Comparative Fit Index (CFI)	0.987	0.967
Tucker-Lewis Index (TLI)	0.981	0.950
Robust Comparative Fit Index (CFI)		NA
Robust Tucker-Lewis Index (TLI)		NA

Root Mean Square Error of Approximation:

RMSEA	0.031	0.045
90 Percent confidence interval - lower	0.000	0.014
90 Percent confidence interval - upper	0.059	0.070
P-value RMSEA <= 0.05	0.847	0.600
Robust RMSEA		NA
90 Percent confidence interval - lower		NA
90 Percent confidence interval - upper		NA

Standardized Root Mean Square Residual:

SRMR	0.083	0.083
------	-------	-------

Parameter Estimates:

Standard errors	Robust.sem
Information	Expected
Information saturated (h1) model	Unstructured

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
visual =~						
x1	1.000				0.639	0.639
x2	0.900	0.188	4.788	0.000	0.575	0.575
x3	0.939	0.197	4.766	0.000	0.600	0.600
textual =~						
x4	1.000				0.835	0.835
x5	0.976	0.118	8.241	0.000	0.815	0.815
x6	1.078	0.125	8.601	0.000	0.900	0.900
speed =~						
x7	1.000				0.471	0.471
x8	1.569	0.461	3.403	0.001	0.740	0.740
x9	1.449	0.409	3.541	0.000	0.683	0.683

Covariances:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
visual ~~						
textual	0.303	0.061	4.981	0.000	0.569	0.569
speed	0.132	0.049	2.700	0.007	0.439	0.439
textual ~~						
speed	0.076	0.046	1.656	0.098	0.192	0.192

Intercepts:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.x1	0.000				0.000	0.000
.x2	0.000				0.000	0.000

.x3	0.000	0.000	0.000
.x4	0.000	0.000	0.000
.x5	0.000	0.000	0.000
.x6	0.000	0.000	0.000
.x7	0.000	0.000	0.000
.x8	0.000	0.000	0.000
.x9	0.000	0.000	0.000
visual	0.000	0.000	0.000
textual	0.000	0.000	0.000
speed	0.000	0.000	0.000

Thresholds:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
x1 t1	-0.388	0.074	-5.223	0.000	-0.388	-0.388
x2 t1	-0.054	0.072	-0.748	0.454	-0.054	-0.054
x3 t1	0.318	0.074	4.309	0.000	0.318	0.318
x4 t1	0.180	0.073	2.473	0.013	0.180	0.180
x5 t1	-0.257	0.073	-3.506	0.000	-0.257	-0.257
x6 t1	1.024	0.088	11.641	0.000	1.024	1.024
x7 t1	0.231	0.073	3.162	0.002	0.231	0.231
x8 t1	1.128	0.092	12.284	0.000	1.128	1.128
x9 t1	0.626	0.078	8.047	0.000	0.626	0.626

Variances:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.x1	0.592				0.592	0.592
.x2	0.670				0.670	0.670
.x3	0.640				0.640	0.640

.x4	0.303				0.303	0.303
.x5	0.336				0.336	0.336
.x6	0.191				0.191	0.191
.x7	0.778				0.778	0.778
.x8	0.453				0.453	0.453
.x9	0.534				0.534	0.534
visual	0.408	0.112	3.651	0.000	1.000	1.000
textual	0.697	0.101	6.883	0.000	1.000	1.000
speed	0.222	0.094	2.363	0.018	1.000	1.000

Scales y*:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
x1	1.000				1.000	1.000
x2	1.000				1.000	1.000
x3	1.000				1.000	1.000
x4	1.000				1.000	1.000
x5	1.000				1.000	1.000
x6	1.000				1.000	1.000
x7	1.000				1.000	1.000
x8	1.000				1.000	1.000
x9	1.000				1.000	1.000

estimated thresholds and tetrachoric correlations

```
> lavInspect (fit, "sampstat")
```

```
$cov
```

	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	1.000								
x2	0.284	1.000							
x3	0.415	0.389	1.000						
x4	0.364	0.328	0.232	1.000					
x5	0.319	0.268	0.138	0.688	1.000				
x6	0.422	0.322	0.206	0.720	0.761	1.000			
x7	-0.048	0.061	0.041	0.200	0.023	-0.029	1.000		
x8	0.159	0.105	0.439	-0.029	-0.059	0.183	0.464	1.000	
x9	0.165	0.210	0.258	0.146	0.183	0.230	0.335	0.403	1.000

```
$mean
```

x1	x2	x3	x4	x5	x6	x7	x8	x9
0	0	0	0	0	0	0	0	0

```
$th
```

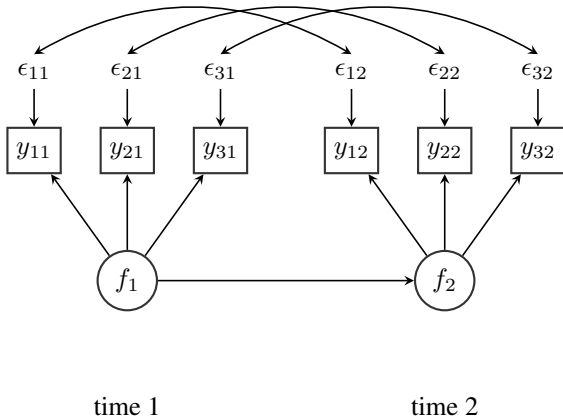
x1 t1	x2 t1	x3 t1	x4 t1	x5 t1	x6 t1	x7 t1	x8 t1	x9 t1
-0.388	-0.054	0.318	0.180	-0.257	1.024	0.231	1.128	0.626

2.11 Panel models for longitudinal data

- panel models postulate *directional* (regression) relationships among the repeated measures
- both within repeated variables (autoregressive) and between repeated variables (cross-lagged)
- focus on the model-implied covariance/correlation structure
- the means are usually ignored
- some subtypes:
 - autoregressive models (the simplex model)
 - cross-lagged models
 - latent autoregressive/cross-lagged models
 - ...

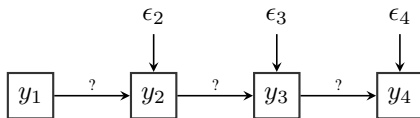
example panel model with a single latent variable

- example with 2 time points:



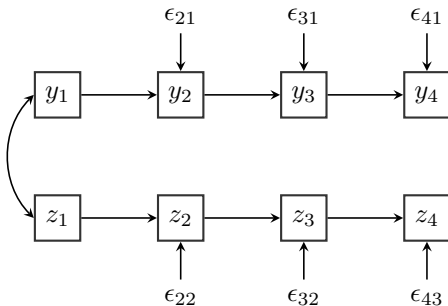
autoregressive models

- each time point is regressed on a previous time point (first order) , or an even further time point (second order, third order, ...)
- alternative names: Markov models, simplex models, panel models, ...
- earliest development dates back to the seminal work of Guttman (1954)
- example first-order univariate autoregressive model:



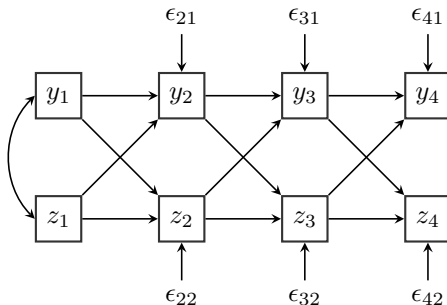
multivariate panel models

- in a multivariate panel model, we have more than one outcome, measured at (the same) t time points
- example: a bivariate panel/simplex model where Y is a measure of mathematical achievement, and Z is a measure of reading ability (4 time points: grade 3, grade 4, grade 5 and grade 6)



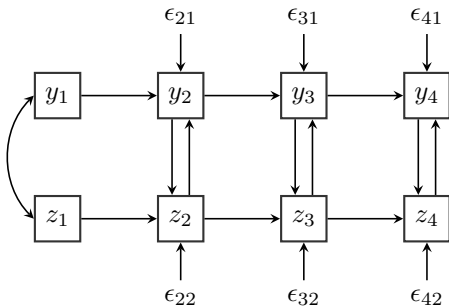
crosslagged effects

- what is the directional effect of one variable on the other?
 - do the two variables develop independently of each other?
 - or does Y exert a greater influence on Z , or vice versa?



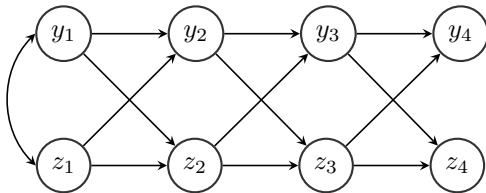
contemporaneous effects

- sometimes, the crossed effects between two variables are not lagged, but contemporaneous (exerting an effect at the same time point)
- this can be unidirectional, or reciprocal
- not everyone believes this approach is useful (in addition: often convergence issues)



panel model with latent variables

- if the ‘repeated’ outcomes are not directly observable, we may replace them with a latent variable with a proper measurement model
- but first, we need to establish ‘measurement invariance’ for the latent variables across time



- in this diagram, the observed indicators have been omitted

2.12 Growth curve models

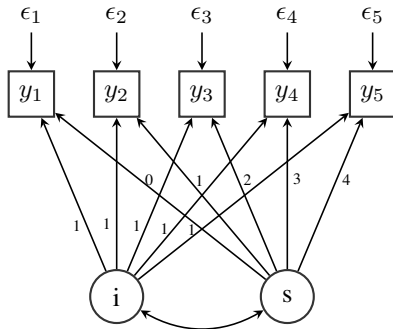
- ‘time’ is typically considered as a continuous variable
- two components:
 - fixed effects: what is the nature of the average trend (linear, quadratic)
 - random effects: individual differences
- in addition, we may try to explain these individual differences by taking into account:
 - time-invariant covariates (age, gender, ...)
 - time-varying covariates (measured at each time point)
- closely related to ‘mixed models’ (linear mixed models, generalized mixed models)
 - limited to balanced data
 - but we can add indirect paths and latent variables
- focus on the mean structure (not the covariance structure)

some references

- Bollen, K.A., & Curran, P.J. (2006). *Latent curve models: A structural equation perspective*. John Wiley & Sons.
- Duncan, T.E., Duncan, S.C., & Strycker, L.A. (2006). *An introduction to latent variable growth curve modeling: Concepts, issues, and applications*. Routledge Academic.
- Preacher, K.J., Wichman, A.L., MacCallum, R.C., & Briggs, N.E. (2008). *Latent Growth Curve Modeling*. Quantitative Applications in the Social Sciences, No. 157, Sage.

a typical growth curve model

- random intercept and random slope



- $y_t = (\text{initial time at time 1}) + (\text{growth per unit time}) \cdot \text{time} + \text{error}$
- $y_t = \text{intercept} + \text{slope} \cdot \text{time} + \text{error}$

example growth curve model: use the lavaan() function

```
> model.syntax <- '  
+ # intercept and slope with fixed coefficients  
+ i =~ 1*t1 + 1*t2 + 1*t3 + 1*t4  
+ s =~ 0*t1 + 1*t2 + 2*t3 + 3*t4  
+  
+ # fixed effects  
+ i ~ 1 # intercept  
+ s ~ 1 # slope  
+  
+ # random effects  
+ i ~~ i # variance random intercept  
+ s ~~ s # variance random slope  
+ i ~~ s # covariance random intercept/slope  
+ '  
> fit <- lavaan(model.syntax, data = Demo.growth, auto.var = TRUE)
```

2.13 Two-level SEM with random intercepts

lavaan syntax setup for two-level SEM

 Σ_B

Between

Within Σ_W

```
model <- '  
  level: 1  
    # here comes the within level  
  level: 2  
    # here comes the between level  
,  
fit <- sem(myModel, myData,  
           cluster = "school")
```

example: Demo.twolevel (simulated data)

- data: 200 clusters, 2500 observations, cluster sizes: 5, 10, 15 and 20
- measures at the within level y_1, y_2, y_3, \dots
- covariates at the within level $x_1, x_2 \dots$
- covariates at the between level w_1 and w_2
- explore the data:

```
> library(lavaan)
> head(round(Demo.twolevel[, c(1:4, 7:12)], 3), n = 9)
```

	y1	y2	y3	y4	x1	x2	x3	w1	w2	cluster
1	0.229	1.356	-0.691	0.803	1.174	-0.623	0.647	-0.248	-0.499	1
2	0.309	-1.862	-2.418	0.766	-1.004	-0.567	0.020	-0.248	-0.499	1
3	0.200	-1.340	0.438	1.197	-0.440	-2.134	-0.459	-0.248	-0.499	1
4	1.045	-0.962	-0.446	-0.203	-0.625	-0.337	1.285	-0.248	-0.499	1
5	0.688	-0.457	-0.642	0.990	-0.845	-0.042	1.560	-0.248	-0.499	1
6	-2.069	-0.600	0.315	0.676	-0.783	-0.224	-0.381	-2.322	-0.691	2
7	-0.787	-0.488	1.132	-0.256	-0.178	-0.583	3.748	-2.322	-0.691	2
8	3.454	1.409	0.930	1.280	0.950	0.259	0.709	-2.322	-0.691	2
9	0.599	-0.291	-1.070	1.930	-1.189	0.815	-0.321	-2.322	-0.691	2

model 1: the empty (univariate) model



y_1

Between

Within

y_1

```
library(lavaan)

model <- '
  level: 1
    y1 ~~ y1
  level: 2
    y1 ~~ y1
  '

fit <- sem(model,
            data = Demo.twolevel,
            cluster = "cluster")

summary(fit, nd = 4)
```

lavaan output (parameter estimates only)

Level 1 []:

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
y1	0.0000			

Variances:

	Estimate	Std.Err	z-value	P(> z)
y1	2.0003	0.0589	33.9574	0.0000

Level 2 []:

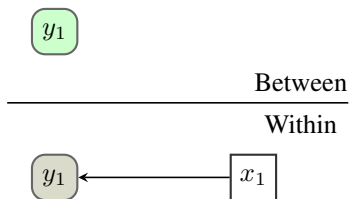
Intercepts:

	Estimate	Std.Err	z-value	P(> z)
y1	0.0198	0.0755	0.2617	0.7935

Variances:

	Estimate	Std.Err	z-value	P(> z)
y1	0.9436	0.1124	8.3931	0.0000

model 2: simple twolevel regression (predictor within)



```
model <- '  
  level: 1  
    y1 ~ x1  
  level: 2  
    y1 ~~ y1  
,  
  
fit <- sem(model,  
           data = Demo.twolevel,  
           cluster = "cluster")  
  
summary(fit, nd = 4)
```

lavaan output (parameter estimates only)

Level 1 []:

Regressions:

	Estimate	Std.Err	z-value	P(> z)
y1 ~ x1	0.4944	0.0276	17.8804	0.0000

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
.y1	0.0000			

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y1	1.7599	0.0518	33.9532	0.0000

Level 2 []:

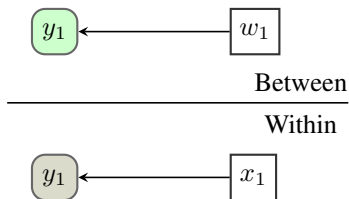
Intercepts:

	Estimate	Std.Err	z-value	P(> z)
.y1	0.0222	0.0745	0.2985	0.7653

Variances:

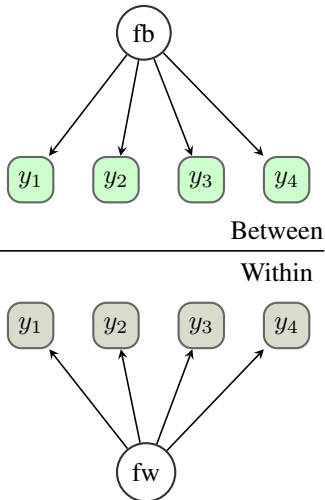
	Estimate	Std.Err	z-value	P(> z)
.y1	0.9367	0.1096	8.5436	0.0000

model 3: simple twolevel regression (no output)



```
model <- '  
  level: 1  
    y1 ~ x1  
  level: 2  
    y1 ~ w1  
,  
  
fit <- sem(model,  
           data = Demo.twolevel,  
           cluster = "cluster")  
  
summary(fit, nd = 4)
```


model 4: one-factor model at both levels



```

model <- '
  level: 1
    fw =~ y1 + y2 + y3 + y4
  level: 2
    fb =~ y1 + y2 + y3 + y4
  ,
fit <- sem(model,
  data = Demo.twolevel,
  cluster = "cluster")

```

lavaan output

```
> summary(fit)
```

```
lavaan 0.6-10 ended normally after 44 iterations
```

Estimator	ML
Optimization method	NLMINB
Number of model parameters	20
Number of observations	2500
Number of clusters [cluster]	200

```
Model Test User Model:
```

Test statistic	1.274
Degrees of freedom	4
P-value (Chi-square)	0.866

```
Parameter Estimates:
```

Standard errors	Standard
Information	Observed
Observed information based on	Hessian

```
Level 1 [within]:
```

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
fw = ~				
y1	1.000			
y2	0.751	0.042	18.051	0.000
y3	0.713	0.040	18.034	0.000
y4	0.315	0.028	11.189	0.000

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
.y1	0.000			
.y2	0.000			
.y3	0.000			
.y4	0.000			
fw	0.000			

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y1	0.949	0.059	15.990	0.000
.y2	1.081	0.044	24.586	0.000
.y3	1.024	0.041	25.177	0.000
.y4	1.080	0.033	32.458	0.000
fw	1.052	0.074	14.269	0.000

Level 2 [cluster]:**Latent Variables:**

	Estimate	Std.Err	z-value	P(> z)
fb =~				
y1	1.000			
y2	0.714	0.056	12.801	0.000
y3	0.579	0.050	11.474	0.000
y4	0.057	0.094	0.611	0.541

Intercepts:

	Estimate	Std.Err	z-value	P(> z)
.y1	0.020	0.076	0.265	0.791
.y2	-0.019	0.061	-0.318	0.750
.y3	-0.045	0.055	-0.817	0.414
.y4	0.022	0.080	0.280	0.779
fb	0.000			

Variances:

	Estimate	Std.Err	z-value	P(> z)
.y1	0.055	0.049	1.122	0.262
.y2	0.122	0.032	3.805	0.000
.y3	0.148	0.028	5.272	0.000
.y4	1.159	0.127	9.111	0.000
fb	0.891	0.122	7.318	0.000

more output

```
> fitMeasures(fit)
```

npar	fmin	chisq	df
20.000	2.904	1.274	4.000
pvalue	baseline.chisq	baseline.df	baseline.pvalue
0.866	1511.382	12.000	0.000
cfi	tli	nnfi	rfi
1.000	1.005	1.005	0.997
nfi	pnfi	ifi	rni
0.999	0.333	1.002	1.002
logl	unrestricted.logl	aic	bic
-16448.595	-16447.958	32937.191	33053.672
ntotal	bic2	rmsea	rmsea.ci.lower
2500.000	32990.127	0.000	0.000
rmsea.ci.upper	rmsea.pvalue	srmr	srmr_within
0.016	1.000	0.020	0.001
srmr_between			
0.018			

```
> lavInspect(fit, "h1")
```

```
$within
$within$cov
  y1  y2  y3  y4
y1 2.000
y2 0.788 1.673
```

```
y3 0.749 0.564 1.557
y4 0.333 0.250 0.231 1.184

$within$mean
  y1    y2    y3    y4
0.001 -0.002 -0.001  0.002

$cluster
$cluster$cov
  y1    y2    y3    y4
y1 0.946
y2 0.635 0.575
y3 0.517 0.368 0.448
y4 0.048 0.019 0.069 1.163

$cluster$mean
  y1    y2    y3    y4
0.019 -0.017 -0.044  0.020

> lavInspect (fit, "implied")

$within
$within$cov
  y1    y2    y3    y4
y1 2.000
y2 0.789 1.673
y3 0.749 0.562 1.558
```

```
y4 0.331 0.248 0.236 1.184
```

```
$within$mean
```

```
y1 y2 y3 y4  
0 0 0 0
```

```
$cluster
```

```
$cluster$cov
```

```
      y1      y2      y3      y4  
y1 0.946  
y2 0.636 0.576  
y3 0.516 0.368 0.447  
y4 0.051 0.036 0.030 1.162
```

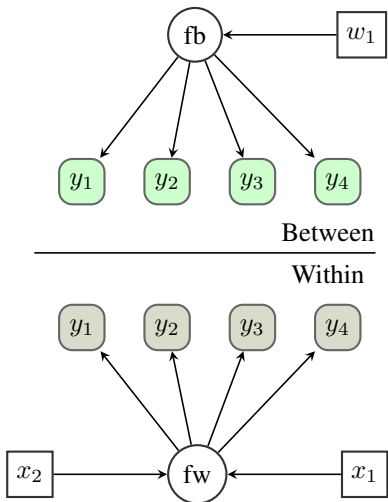
```
$cluster$mean
```

```
      y1      y2      y3      y4  
0.020 -0.019 -0.045 0.022
```

```
> lavInspect (fit, "icc")
```

```
      y1      y2      y3      y4  
0.321 0.256 0.223 0.495
```

model 5: adding covariates (no output)



```

model <- '
  level: 1
    fw =~ y1 + y2 + y3 + y4
    fw ~ x1 + x2

  level: 2
    fb =~ y1 + y2 + y3 + y4
    fb ~ w1
'

fit <- sem(model,
  data = Demo.twolevel,
  cluster = "cluster")

```


...and our time is up.

Thank you for following this workshop.

Thank you for choosing open-source software.