

Psychology 454: Latent Variable Modeling

further adventures with lavaan

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February, 2011

Outline

- 1 Introduction to CFA/SEM programs
 - What is lavaan?
 - lavaan syntax
- 2 Confirmatory Factor Analysis
 - A simple confirmatory analysis
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 - Fixing parameters - starting values and equality constraints
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 - Means structure
 - Multiple groups
 - Measurement invariance
 - Growth Curve analysis
 - Modifying the model
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Latent Variable Modeling programs

- Commercial programs
 - LISREL/PRELIS (Jöreskog, 1978; Joreskog & Sorbom, 1993; Jöreskog & Sörbom, 1999) <http://www.ssicentral.com/lisrel/techdocs/IPUG.pdf> Users Manual
 - EQS (Bentler, 1995)
 - AMOS Arbuckle (1989, 1994) [http://spss.wikia.com/wiki/SEM_\(structural_equation_modeling\)-_Amos](http://spss.wikia.com/wiki/SEM_(structural_equation_modeling)-_Amos) Amos wiki
 - MPLUS (Muthén & Muthén, 2007) <http://www.statmodel.com/ugexcerpts.shtml> User's guide with examples
- Open source
 - Mx (Neale, 1994)
 - OpenMx
 - sem (Fox, 2009)
 - lavaan (Rosseel, 2010)

- The lavaan package is free open-source software. This means (among other things) that there is no warranty whatsoever.
- The numerical results of the lavaan package are typically very close, if not identical, to the results of the commercial package Mplus. If you wish to compare the results with other SEM packages, you can use the optional argument `mimic="EQS"` when calling the `cfa`, `sem` or `growth` functions
- The lavaan package is not finished yet. But it is already very useful for most users, or so we hope. There are a number of known minor issues and some features are simply not implemented yet.
- Some important features that are currently not available in lavaan are:
 - support for categorical/censored variables
 - support for discrete latent variables (mixture models)
 - support for hierarchical/multilevel datasets

$$y \sim f1 + f2 + x1 + x2$$
$$f_1 \sim f_2 + f_3$$
$$f_2 \sim f_3 + x_1 + x_2$$

Latent variables

$$f1 \approx y1 + y2 + y3$$
$$f_2 = y_4 + y_5 + y_6$$
$$f_3 = y_7 + y_8 + y_9 + y_{10}$$

Variances and covariances

$$y1 \sim y1$$
$$y1 \sim y2$$
$$f1 \sim f2$$

Intercepts

$$y_1 \sim 1$$
$$f1 \sim 1$$

Entering the model syntax as a string literal

```
myModel <- ' # regressions
y1 + y2 ~ f1 + f2 + x1 + x2
f1 ~ f2 + f3
f2 ~ f3 + x1 + x2
# latent variable definitions
f1 =~ y1 + y2 + y3
f2 =~ y4 + y5 + y6
f3 =~ y7 + y8 +
y9 + y10
# variances and covariances
y1 ~~ y1
y1 ~~ y2
f1 ~~ f2
# intercepts
y1 ~ 1
f1 ~ 1
'
```

The HolzingerSwineford data set

```
HS.model <- '  
visual =~ x1 + x2 + x3  
  textual =~ x4 + x5 + x6  
  speed =~ x7 + x8 + x9  
'  
  
fit <- cfa(HS.model, data = HolzingerSwineford1939)  
  
summary(fit, fit.measures = TRUE)  
lavaan.diagram(fit) #need to use the newer function
```


A simple confirmatory analysis

Holzinger Swineford analysis

Lavaan (0.4-7) converged normally after 35 iterations Root Mean Square Error of Approximation:

Number of observations	301	RMSEA		0.092
		90 Percent Confidence Interval	0.071	0.114
Estimator	ML	P-value RMSEA <= 0.05		0.001
Minimum Function Chi-square	85.306			
Degrees of freedom	24	Standardized Root Mean Square Residual:		
P-value	0.000			
		SRMR		0.065

Chi-square test baseline model:

Parameter estimates:				
Minimum Function Chi-square	918.852			
Degrees of freedom	36	Information		Expected
P-value	0.000	Standard Errors		Standard

Full model versus baseline model:

		Estimate	Std.err	Z-value	P(> z)	
Latent variables:						
Comparative Fit Index (CFI)	0.931	visual =~				
Tucker-Lewis Index (TLI)	0.896	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

Loglikelihood and Information Criteria:

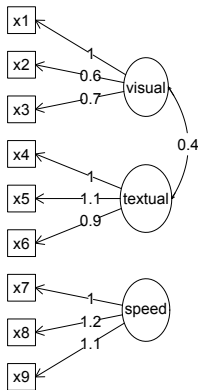
Loglikelihood user model (H0)	-3737.745
Loglikelihood unrestricted model (H1)	-3695.092
Number of free parameters	21
Akaike (AIC)	7517.490
Bayesian (BIC)	7595.339
Sample-size adjusted Bayesian (BIC)	7528.739

more parameters

					Variances:				
					x1				
					x2	0.549	0.114	4.833	0.000
					x3	1.134	0.102	11.146	0.000
					x4	0.844	0.091	9.317	0.000
					x5	0.371	0.048	7.778	0.000
Covariances:	visual ~~				x6	0.446	0.058	7.642	0.000
	textual	0.408	0.074	5.552	0.000	x7	0.356	0.043	8.277
	speed	0.262	0.056	4.660	0.000	x8	0.799	0.081	9.823
	textual ~~					x9	0.488	0.074	6.573
	speed	0.173	0.049	3.518	0.000	visual	0.566	0.071	8.003
					textual	0.809	0.145	5.564	0.000
					speed	0.979	0.112	8.737	0.000
						0.384	0.086	4.451	0.000

Graphic output using (revised) lavaan.diagram

Confirmatory structure



Redo with alternative parameterization

```

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

fit <- cfa(HS.model, data = HolzingerSwineford1939, std.ov=TRUE, std.lv=TRUE)

summary(fit, fit.measures = TRUE)
lavaan.diagram(fit) #need to use the newer function

```

	RMSEA	0.092
--	-------	-------

Number of observations	301	AKISER		01002
		99 Percent Confidence Interval	0.071	0.114

90 Percent Confidence Interval 0.071 0.114

Estimator	MI	P-value RMSEA <= 0.05	0.001
-----------	----	-----------------------	-------

Estimator	ML
Model 1: $\beta_1 = 0.000$	0.000

Minimum Function Chi-square 85.306 Standardized Root Mean Square Residual:

Degrees of freedom 24 Standardized Root Mean Square Residual:

P-value 0.000

Parameter estimates:

Minimum Function Chi-square 918.852

Minimum function on square	518.882	Information	Expected
----------------------------	---------	-------------	----------

Degrees of freedom	36	Standard Errors	Standard
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P-value	0.000	Standard Error	Standard Error
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Latent variables:

visual = ~

Comparative Fit Index (CFI) 0.931 visual =

Comparative Fit Index (CFI)	0.951	x1	0.771	0.069	11.127	0.000
Tucker-Lewis Index (TLI)	0.906					

Lucker-Lewis Index (LLI)	0.896	x2	0.423	0.066	6.429	0.000
--------------------------	-------	----	-------	-------	-------	-------

x2	0.580	0.066	8.817	0.000
----	-------	-------	-------	-------

textual =

textual = ~

										x4	0.850	0.049	17.474	0.000
--	--	--	--	--	--	--	--	--	--	----	-------	-------	--------	-------

Loglikelihood user model (H0)	-3422.624	x5	0.854	0.049	17.576	0.000
-------------------------------	-----------	----	-------	-------	--------	-------

Loglikelihood unrestricted model (H1)	-3379.971	0.004	0.049	17.910	0.000
---------------------------------------	-----------	-------	-------	--------	-------

x6	0.837	0.049	17.082	0.000
----	-------	-------	--------	-------

Number of free parameters 21 speed = τ

Number of free parameters	21	x7	0.569	0.064	8.903	0.000
---------------------------	----	----	-------	-------	-------	-------

Akaike (AIC)	6887.248	x7	0.000	0.000	0.000	0.000
		x8	0.722	0.065	11.090	0.000

Bayesian (BIC)	6965.097	0.722	0.085	11.090	0.000
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Sample-size adjusted Bayesian (BIC)	6898.497	x9	0.664	0.064	10.305	0.000
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sample-size adjusted Bayesian (DIC)	6696.497
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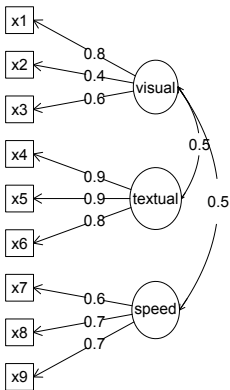
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A simple confirmatory analysis

					Variances:					
					x1	0.403	0.083	4.833	0.000	
					x2	0.818	0.073	11.146	0.000	
					x3	0.660	0.071	9.317	0.000	
					x4	0.274	0.035	7.779	0.000	
Covariances:	visual ~~				x5	0.268	0.035	7.642	0.000	
	textual	0.459	0.064	7.189	0.000	x6	0.297	0.036	8.277	0.000
	speed	0.471	0.073	6.461	0.000	x7	0.673	0.069	9.823	0.000
	textual ~~				x8	0.476	0.072	6.573	0.000	
	speed	0.283	0.069	4.117	0.000	x9	0.556	0.069	8.003	0.000
					visual	1.000				
					textual	1.000				
					speed	1.000				

The standardized solution to the Holzinger Swineford 1939 problem

Confirmatory structure



EFA of the Holzinger Swineford 1939 problem

```
f3 <- fa(HolzingerSwineford1939[7:15],3)
f3
diagram(f3,cut=.1)
```

```
Factor Analysis using method = minres
Call: fa(r = HolzingerSwineford1939[7:15],
      nfactors = 3)
```

```
Standardized loadings based upon
correlation matrix
```

	MR1	MR3	MR2	h2	u2
x1	0.19	0.60	0.03	0.49	0.51
x2	0.04	0.51	-0.12	0.25	0.75
x3	-0.07	0.69	0.02	0.46	0.54
x4	0.84	0.02	0.01	0.72	0.28
x5	0.89	-0.07	0.01	0.76	0.24
x6	0.81	0.08	-0.01	0.69	0.31
x7	0.04	-0.15	0.72	0.50	0.50
x8	-0.03	0.10	0.70	0.53	0.47
x9	0.03	0.37	0.46	0.46	0.54

	MR1	MR3	MR2
SS loadings	2.24	1.34	1.28
Proportion Var	0.25	0.15	0.14
Cumulative Var	0.25	0.40	0.54

```
With factor correlations of
MR1 MR3 MR2
MR1 1.00 0.33 0.22
MR3 0.33 1.00 0.27
MR2 0.22 0.27 1.00
```

Test of the hypothesis that 3 factors are sufficient.

The degrees of freedom for the null model are 36 and the objective function was 3.05 with Chi Square of 904.1
The degrees of freedom for the model are 12 and the objective function was 0.08

The root mean square of the residuals is 0.01
The df corrected root mean square of the residuals is 0.03
The number of observations was 301 with
Chi Square = 22.38 with prob < 0.034

Tucker Lewis Index of factoring reliability = 0.964
RMSEA index = 0.055 and the 90 % confidence intervals are
0.054 0.06

BIC = -46.11
Fit based upon off diagonal values = 1
Measures of factor score adequacy

	MR1	MR3	MR2
Correlation of scores with factors	0.94	0.84	0.85
Multiple R square of scores with factors	0.89	0.71	0.72
Minimum correlation of possible factor scores	0.78	0.42	0.45

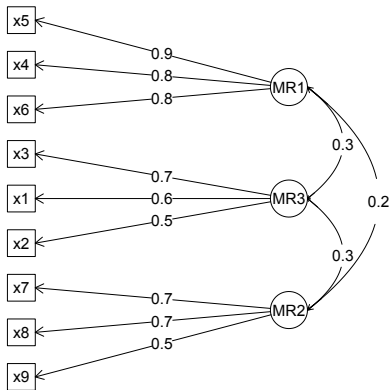
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compare with EFA

Holzing Swineford EFA – compare with CFA

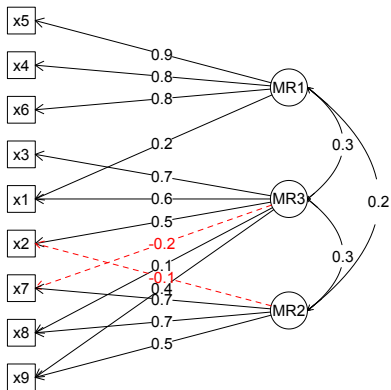
Factor Analysis



Holzinger Swineford EFA – simple is FALSE

```
diagram(f3,cut=.1,simple=FALSE)
```

Factor Analysis



Fixing parameters to simply models

- Perhaps the greatest power of SEM type programs is the ability to fix parameters or to force equality constraints.
 - EFA allows all parameters to vary
 - CFA allows only certain parameters to vary
- Can fix covariances to be zero
- Can fix different paths to be equal

Two ways of fixing the covariances to be 0

```
HS.ortho <- '
# three-factor model
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ NA*x7 + x8 + x9
# orthogonal factors
visual ~~ 0*speed
textual ~~ 0*speed
# fix variance of speed factor
speed ~~ 1*speed'
fit.hs.ortho <- cfa(HS.ortho, data=HolzingerSwineford1939, std.ov=TRUE, std.lv=TRUE)
```

or (if they are all to be zero)

```
HS.model <- ' visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9 '
fit.HS.ortho <- cfa(HS.model, data=HolzingerSwineford1939, orthogonal=TRUE)
```

Lavaan (0.4-7) converged normally after 20 iterations

Number of observations	301
Estimator	ML
Minimum Function Chi-square	117.946
Degrees of freedom	26
P-value	0.000

Fixing parameters - starting values and equality constraints

Not as good a fit

```
> summary(fit.hs.ortho,fit.measures=TRUE)
```

Lavaan (0.4-7) converged normally after 20 iterations

Root Mean Square Error of Approximation:

Number of observations	301	RMSEA	0.108
Estimator	ML	90 Percent Confidence Interval	0.089 0.129
Minimum Function Chi-square	117.946	P-value RMSEA <= 0.05	0.000
Degrees of freedom	26	Standardized Root Mean Square Residual:	
P-value	0.000	SRMR	0.125

Chi-square test baseline model:

Parameter estimates:

Minimum Function Chi-square	918.852	Information	Expected
Degrees of freedom	36	Standard Errors	Standard
P-value	0.000		

Full model versus baseline model:

Latent variables:

		Estimate	Std.err	Z-value	P(> z)
Comparative Fit Index (CFI)	0.896				
Tucker-Lewis Index (TLI)	0.856				

Loglikelihood and Information Criteria:

Loglikelihood user model (H0)	-3438.944	visual =~			
Loglikelihood unrestricted model (H1)	-3379.971	x1	0.777	0.075	10.376 0.000
		x2	0.430	0.067	6.423 0.000
		x3	0.568	0.069	8.270 0.000
		textual =~			
		x4	0.851	0.049	17.491 0.000
		x5	0.853	0.049	17.545 0.000
		x6	0.837	0.049	17.079 0.000
		speed =~			
Number of free parameters	19	x7	0.607	0.067	9.040 0.000
Akaike (AIC)	6915.889	x8	0.800	0.073	10.899 0.000
Bayesian (BIC)	6986.324	x9	0.560	0.066	8.509 21.05000
Sample-size adjusted Bayesian (BIC)	6926.067				

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Fixing parameters - starting values and equality constraints

					Variances:				
					speed	1.000			
					x1	0.394	0.095	4.155	0.000
					x2	0.811	0.074	10.965	0.000
					x3	0.675	0.074	9.085	0.000
					x4	0.273	0.035	7.735	0.000
					x5	0.269	0.035	7.662	0.000
					x6	0.297	0.036	8.263	0.000
					x7	0.629	0.073	8.650	0.000
					x8	0.357	0.094	3.794	0.000
					x9	0.683	0.071	9.640	0.000
					visual	1.000			
					textual	1.000			
Covariances:									
visual	~~								
speed		0.000							
textual	~~								
speed		0.000							
visual	~~								
textual		0.461	0.064	7.195	0.000				

Fixing starting values

(If you have a problem with a solution, you can help it if you give it a reasonable starting location.)

```
visual =~ x1 + start(0.8)*x2 + start(1.2)*x3  
textual =~ x4 + start(0.5)*x5 + start(1.0)*x6  
speed =~ x7 + start(0.7)*x8 + start(1.8)*x9
```

This technique works for all SEM programs (although details vary). The reason to give good starting values is that the search optimization can get bogged down in the wrong part of the parameter space.

Automatic naming

```
> model <- '
+ # latent variable definitions
+ ind60 =~ x1 + x2 + x3
+ dem60 =~ y1 + y2 + y3 + y4
+ dem65 =~ y5 + y6 + y7 + y8
+ # regressions
+ dem60 ~ ind60
+ dem65 ~ ind60 + dem60
+ # residual (co)variances
+ y1 ~~ y5
+ y2 ~~ y4 + y6
+ y3 ~~ y7
+ y4 ~~ y8
+ y6 ~~ y8
+ '
> fit <- sem(model, data=PoliticalDemocracy)
coef(fit)

ind60=~x2 ind60=~x3 dem60=~y2 dem60=~y3 dem60=~y4 dem65=~y6
2.180 1.819 1.257 1.058 1.265 1.186
dem65=~y7 dem65=~y8 dem60~ind60 dem65~ind60 dem65~dem60 y1~~y5
1.280 1.266 1.483 0.572 0.837 0.624
y2~~y4 y2~~y6 y3~~y7 y4~~y8 y6~~y8 x1~~x1
1.313 2.153 0.795 0.348 1.356 0.082
x2~~x2 x3~~x3 y1~~y1 y2~~y2 y3~~y3 y4~~y4
0.120 0.467 1.891 7.373 5.068 3.148
y5~~y5 y6~~y6 y7~~y7 y8~~y8 ind60~~ind60 dem60~~dem60
2.351 4.954 3.431 3.254 0.448 3.956
dem65~~dem65
0.172
```


Specifying the name

```
> model <- '
+ # latent variable definitions
+ ind60 =~ x1 + x2 + label("myLabel")*x3
+ dem60 =~ y1 + y2 + y3 + y4
+ dem65 =~ y5 + y6 + y7 + y8
+ # regressions
+ dem60 ~ ind60
+ dem65 ~ ind60 + dem60
+ # residual (co)variances
+ y1 ~~ y5
+ y2 ~~ y4 + y6
+ y3 ~~ y7
+ y4 ~~ y8
+ y6 ~~ y8
+ '
```

Using names to specify equality constraints

```
visual =~ x1 + x2 + equal("visual=~x2")*x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
```

Estimate the means

```

HS.model.means <- '
# three-factor model
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
# intercepts
x1 ~ 1
x2 ~ 1
x3 ~ 1
x4 ~ 1
x5 ~ 1
x6 ~ 1
x7 ~ 1
x8 ~ 1
x9 ~ 1'
fit.means <- cfa(HS.model.means, data = HolzingerSwineford1939, std.lv=
TRUE)

or
fit <- cfa(HS.model, data = HolzingerSwineford1939, meanstructure = TRUE)
summary(fit.means, fit.measures=TRUE)

```

Just show the parameter estimates

	Estimate	Std.err	Z-value	P(> z)	Intercepts:				
					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
Latent variables:					x4	3.061	0.067	45.694	0.000
visual =~					x5	4.341	0.074	58.452	0.000
x1	0.900	0.081	11.127	0.000	x6	2.186	0.063	34.667	0.000
x2	0.498	0.077	6.429	0.000	x7	4.186	0.063	66.766	0.000
x3	0.656	0.074	8.817	0.000	x8	5.527	0.058	94.854	0.000
textual =~					x9	5.374	0.058	92.546	0.000
x4	0.990	0.057	17.474	0.000	visual	0.000			
x5	1.102	0.063	17.576	0.000	textual	0.000			
x6	0.917	0.054	17.082	0.000	speed	0.000			
speed =~									
x7	0.619	0.070	8.903	0.000					
x8	0.731	0.066	11.090	0.000	Variances:				
x9	0.670	0.065	10.305	0.000	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
Covariances:					x4	0.371	0.048	7.779	0.000
visual ~~					x5	0.446	0.058	7.642	0.000
textual	0.459	0.064	7.189	0.000	x6	0.356	0.043	8.277	0.000
speed	0.471	0.073	6.461	0.000	x7	0.799	0.081	9.823	0.000
textual ~~					x8	0.488	0.074	6.573	0.000
speed	0.283	0.069	4.117	0.000	x9	0.566	0.071	8.003	0.000
					visual	1.000			
					textual	1.000			
					speed	1.000			

More useful if we want to fix some intercepts to be different from others

```
# three-factor model
visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9
# intercepts with fixed values
x1 ~ 0.5*1
x2 ~ 0.5*1
x3 ~ 0.5*1
x4 ~ 0.5*1
```

Analyzing multiple groups

- When studying differences in ages, gender, school, it is useful to be able to model them separately, but to get an overall goodness of fit.
 - Does a basic structure hold in different groups?
- More importantly, we can ask if the parameters in the two groups are the same. That is, we can add equality constraints.
- We can examine equality of the loadings, equality of the covariances, equality of the mean structure.

Multiple groups

```

HS.model <- ' visual =~ x1 + x2 + x3
textual =~ x4 + x5 + x6
speed =~ x7 + x8 + x9 '
fit <- cfa(HS.model, data=HolzingerSwineford1939, group="school")
summary(fit)

```

Lavaan (0.4-7) converged normally after 60 iterations

Number of observations per group	
Pasteur	156
Grant-White	145

Estimator	ML
Minimum Function Chi-square	115.851
Degrees of freedom	48
P-value	0.000

Chi-square for each group:

Pasteur	64.309
Grant-White	51.542

Parameter estimates:

Information	Expected
Standard Errors	Standard

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Multiple groups

Group 1 [Pasteur]:

Group 2 [Grant-White]:

	Estimate	Std.err	Z-value	P(> z)		Estimate	Std.err	Z-value	P(> z)
Latent variables:					Latent variables:				
visual =~					visual =~				
x1	1.000				x1	1.000			
x2	0.394	0.122	3.220	0.001	x2	0.736	0.155	4.760	0.000
x3	0.570	0.140	4.076	0.000	x3	0.925	0.166	5.584	0.000
textual =~					textual =~				
x4	1.000				x4	1.000			
x5	1.183	0.102	11.613	0.000	x5	0.990	0.087	11.418	0.000
x6	0.875	0.077	11.421	0.000	x6	0.963	0.085	11.377	0.000
speed =~					speed =~				
x7	1.000				x7	1.000			
x8	1.125	0.277	4.057	0.000	x8	1.226	0.187	6.569	0.000
x9	0.922	0.225	4.104	0.000	x9	1.058	0.165	6.429	0.000
Covariances:					Covariances:				
visual ~~					visual ~~				
textual	0.479	0.106	4.531	0.000	textual	0.408	0.098	4.153	0.000
speed	0.185	0.077	2.397	0.017	speed	0.276	0.076	3.639	0.000
textual ~~					textual ~~				
speed	0.182	0.069	2.628	0.009	speed	0.222	0.073	3.022	0.003
Variances:					Variances:				
x1	0.298	0.232	1.286	0.198	x1	0.715	0.126	5.675	0.000
x2	1.334	0.158	8.464	0.000	x2	0.899	0.123	7.339	0.000
x3	0.989	0.136	7.271	0.000	x3	0.557	0.103	5.409	0.000
x4	0.425	0.069	6.138	0.000	x4	0.315	0.065	4.870	0.000
x5	0.456	0.086	5.292	0.000	x5	0.419	0.072	5.812	0.000
x6	0.290	0.050	5.780	0.000	x6	0.406	0.069	5.880	0.000
x7	0.820	0.125	6.580	0.000	x7	0.600	0.091	6.584	0.000
x8	0.510	0.116	4.406	0.000	x8	0.401	0.094	4.248	0.000
x9	0.680	0.104	6.516	0.000	x9	0.535	0.089	6.010	0.000
visual	1.000	0.276	3.667	0.000	visual	0.733	0.122	6.010	0.000

Multiple groups, multiple constraints

- Can constrain a single parameter to be equal across groups
 - Use the naming convention and the equal comand
- Can constrain equivalent parameters across groups to be equal (group.equal)

Equal parameters across groups

```
HS.model <- ' visual =~ x1 + x2 + x3
  textual =~ x4 + x5 + x6
  speed =~ x7 + x8 + x9 '
fit <- cfa(HS.model, data=HolzingerSwineford1939, group="school",
  group.equal=c("loadings"),std.ov=TRUE,std.lv=TRUE)
summary(fit)
```

Lavaan (0.4-7) converged normally after 27 iterations

Number of observations per group

Pasteur	156
Grant-White	145

Estimator	ML
Minimum Function Chi-square	122.862
Degrees of freedom	57
P-value	0.000

Chi-square for each group:

Pasteur	68.598
Grant-White	54.264

35 0.0
56 0.0

Group 1 [Pasteur]:					Group 2 [Grant-White]:				
	Estimate	Std.err	Z-value	P(> z)		Estimate	Std.err	Z-value	P(> z)
Latent variables:					Latent variables:				
visual =~					visual =~				
x1	0.737	0.066	11.100	0.000	x1	0.737	0.066	11.100	0.000
x2	0.448	0.065	6.888	0.000	x2	0.448	0.065	6.888	0.000
x3	0.625	0.065	9.658	0.000	x3	0.625	0.065	9.658	0.000
textual =~					textual =~				
x4	0.841	0.049	17.127	0.000	x4	0.841	0.049	17.127	0.000
x5	0.843	0.049	17.207	0.000	x5	0.843	0.049	17.207	0.000
x6	0.830	0.049	16.826	0.000	x6	0.830	0.049	16.826	0.000
speed =~					speed =~				
x7	0.597	0.063	9.490	0.000	x7	0.597	0.063	9.490	0.000
x8	0.736	0.064	11.559	0.000	x8	0.736	0.064	11.559	0.000
x9	0.644	0.063	10.221	0.000	x9	0.644	0.063	10.221	0.000
Covariances:					Covariances:				
visual ~~					visual ~~				
textual	0.467	0.087	5.369	0.000	textual	0.537	0.085	6.315	0.000
speed	0.343	0.109	3.149	0.002	speed	0.530	0.097	5.477	0.000
textual ~~					textual ~~				
speed	0.333	0.094	3.527	0.000	speed	0.331	0.093	3.557	0.000
Variances:					Variances:				
x1	0.421	0.095	4.438	0.000	x1	0.479	0.095	5.043	0.000
x2	0.823	0.102	8.040	0.000	x2	0.759	0.098	7.758	0.000
x3	0.634	0.096	6.635	0.000	x3	0.567	0.089	6.399	0.000
x4	0.316	0.052	6.137	0.000	x4	0.260	0.048	5.399	0.000
x5	0.267	0.048	5.590	0.000	x5	0.302	0.052	5.817	0.000
x6	0.299	0.050	6.042	0.000	x6	0.312	0.052	5.998	0.000
x7	0.698	0.098	7.095	0.000	x7	0.577	0.084	6.894	0.000
x8	0.528	0.100	5.303	0.000	x8	0.379	0.081	4.703	0.000

Do the measures measure the same construct across groups?

- Is the configuration the same?
 - Most abstract level of invariance – “are the arrows the same”
- Weak invariance – are the loadings the same?
- Strong invariance – equal loadings + intercepts

Testing for measurement invariance

```
measurementInvariance(HS.model, data = HolzingerSwineford1939,
  group = "school")
```

Measurement invariance tests:

Model 1: configural invariance:

chisq	df	pvalue	cfi	rmsea	bic
115.851	48.000	0.000	0.923	0.097	7604.094

Model 2: weak invariance (equal loadings):

chisq	df	pvalue	cfi	rmsea	bic
124.044	54.000	0.000	0.921	0.093	7578.043

[Model 1 versus model 2]

delta.chisq	delta.df	delta.p.value	delta.cfi
8.192	6.000	0.224	0.002

Model 3: strong invariance (equal loadings + intercepts):

chisq	df	pvalue	cfi	rmsea	bic
164.103	60.000	0.000	0.882	0.107	7686.588

[Model 1 versus model 3]

delta.chisq	delta.df	delta.p.value	delta.cfi
48.251	12.000	0.000	0.041

[Model 2 versus model 3]

delta.chisq	delta.df	delta.p.value	delta.cfi
40.059	6.000	0.000	0.038

Model 4: equal loadings + intercepts + means:

chisq	df	pvalue	cfi	rmsea	bic
204.605	63.000	0.000	0.854	0.122	7709.969

Demo.growth : A toy data set

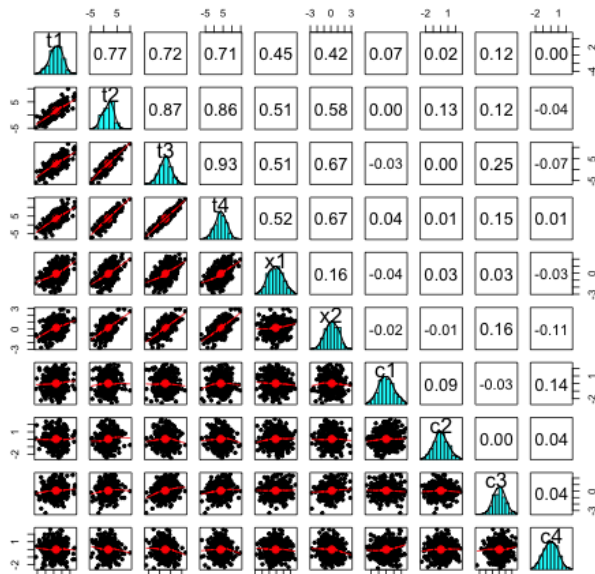
```
data(Demo.growth)
describe(Demo.growth)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	skew	kurtosis	se
t1	1	400	0.59	1.58	0.67	0.64	1.50	-4.35	5.21	9.56	-0.18	0.23	0.08
t2	2	400	1.67	2.13	1.88	1.69	2.10	-4.86	9.95	14.81	-0.02	0.48	0.11
t3	3	400	2.59	2.72	2.73	2.62	2.62	-6.02	11.53	17.55	-0.14	0.46	0.14
t4	4	400	3.64	3.38	3.72	3.69	3.14	-7.34	14.72	22.06	-0.13	0.44	0.17
x1	5	400	-0.09	1.03	-0.08	-0.11	1.11	-2.82	2.72	5.54	0.08	-0.33	0.05
x2	6	400	0.14	0.96	0.13	0.15	1.01	-2.83	2.88	5.71	-0.12	0.01	0.05
c1	7	400	0.01	0.99	-0.03	-0.01	0.98	-2.58	2.57	5.14	0.11	-0.28	0.05
c2	8	400	0.03	0.95	0.01	0.02	0.87	-2.54	2.71	5.25	0.13	-0.07	0.05
c3	9	400	0.07	0.93	0.07	0.06	0.93	-3.40	2.61	6.01	-0.02	0.38	0.05
c4	10	400	-0.02	0.92	-0.01	-0.02	0.95	-2.45	2.65	5.11	0.02	-0.21	0.05

Data.growth: A toy data set

- ① t1 Measured value at time point 1
- ② t2 Measured value at time point 2
- ③ t3 Measured value at time point 3
- ④ t4 Measured value at time point 4
- ⑤ x1 Predictor 1 influencing intercept and slope
- ⑥ x2 Predictor 2 influencing intercept and slope
- ⑦ c1 Time-varying covariate time point 1
- ⑧ c2 Time-varying covariate time point 2
- ⑨ c3 Time-varying covariate time point 3
- ⑩ c4 Time-varying covariate time point 4

Growth: SPLOM



Fitting a growth model to the toy problem

```
model <- ' i =~ 1*t1 + 1*t2 + 1*t3 + 1*t4
s =~ 0*t1 + 1*t2 + 2*t3 + 3*t4 '
fit <- growth(model, data=Demo.growth)
summary(fit)
```

Lavaan (0.4-7) converged normally after 43 iterations

Number of observations	400
Estimator	ML
Minimum Function Chi-square	8.069
Degrees of freedom	5
P-value	0.152

Parameter estimates:

Information	Expected
Standard Errors	Standard

Estimate	Std.err	Z-value	P(> z)
----------	---------	---------	---------

Growth model with parameter values

Latent variables:

$$i = z$$

t1	1.000
t2	1.000
t3	1.000
t4	1.000

$$S = 2$$

t1	0.000
t2	1.000
t3	2.000
t4	3.000

Covariances:

i ~ ~

s	0.618	0.071	8.686	0.000
---	-------	-------	-------	-------

Intercepts:

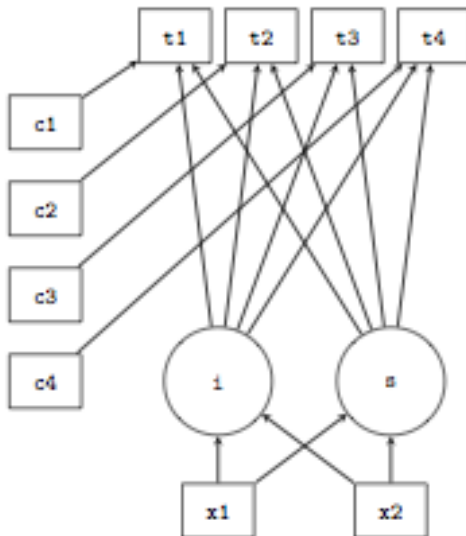
t1	0.000
t2	0.000
t3	0.000
t4	0.000

i	0.615	0.077	8.007	0.000
s	1.006	0.042	24.076	0.000

Variances:

t1	0.595	0.086	6.944	0.000
t2	0.676	0.061	11.061	0.000
t3	0.635	0.072	8.761	0.000
t4	0.508	0.124	4.090	0.000
i	1.932	0.173	11.194	0.000
s	0.587	0.052	11.336	0.000

- “Technically, the growth function is almost identical to the sem function. But a meanstructure is automatically assumed, and the observed intercepts are fixed to zero by default, while the latent variable intercepts and means are freely estimated.
- A slightly more complex model adds two regressors (x_1 and x_2) that influence the latent growth factors.
- In addition, a time-varying covariate that influences the outcome measure at the four time points has been added to the model.
- A graphical representation of this model together with the corresponding lavaan syntax is presented”.



Linear growth with time-varying covariates

```
# a linear growth model with a time-varying covariate
model <- '
# intercept and slope with fixed coefficients
i =~ 1*t1 + 1*t2 + 1*t3 + 1*t4
s =~ 0*t1 + 1*t2 + 2*t3 + 3*t4
# regressions
i ~ x1 + x2
s ~ x1 + x2
# time-varying covariates
t1 ~ c1
t2 ~ c2
t3 ~ c3
t4 ~ c4
'

fit <- growth(model, data=Demo.growth)
summary(fit)
```

Lavaan (0.4-7) converged normally after 40 iterations

Number of observations	400
Estimator	ML
Minimum Function Chi-square	26.059
Degrees of freedom	21

Parameters of the growth model

	Estimate	Std.err	Z-value	P(> z)					
Latent variables:									
i =~									
t1	1.000								
t2	1.000								
t3	1.000				Covariances:				
t4	1.000				i ~~				
s =~					s	0.075	0.040	1.855	0.064
t1	0.000								
t2	1.000				Intercepts:				
t3	2.000				t1	0.000			
t4	3.000				t2	0.000			
					t3	0.000			
					t4	0.000			
Regressions:					i	0.580	0.062	9.368	0.000
i ~					s	0.958	0.029	32.552	0.000
x1	0.608	0.060	10.134	0.000					
x2	0.604	0.064	9.412	0.000	Variances:				
s ~					t1	0.580	0.080	7.230	0.000
x1	0.262	0.029	9.198	0.000	t2	0.596	0.054	10.969	0.000
x2	0.522	0.031	17.083	0.000	t3	0.481	0.055	8.745	0.000
t1 ~					t4	0.535	0.098	5.466	0.000
c1	0.143	0.050	2.883	0.004	i	1.079	0.112	9.609	0.000
t2 ~					s	0.224	0.027	8.429	0.000
c2	0.289	0.046	6.295	0.000					
t3 ~									
c3	0.328	0.044	7.361	0.000					
t4 ~									
c4	0.330	0.058	5.655	0.000					

Modifying a model

- There are many reasons a model does not fit.
 - In particular, some paths may be badly fit (usually because they were ignored).
 - How much will a parameter change (Expected Parameter Change) if a parameter is adjusted.

Modification indices for the HS problem

```
fit <- cfa(HS.model, data = HolzingerSwineford1939)
mi <- modindices(fit)
mi[mi$op == "=", ]
```

	lhs	op	rhs	mi	epc	sepc.lv	sepc.all								
1	visual	=~	x1	NA	NA	NA	NA	14	textual	=~	x5	0.000	0.000	0.000	0.000
2	visual	=~	x2	0.000	0.000	0.000	0.000	15	textual	=~	x6	0.000	0.000	0.000	0.000
3	visual	=~	x3	0.000	0.000	0.000	0.000	16	textual	=~	x7	0.098	-0.021	-0.021	-0.019
4	visual	=~	x4	1.211	0.077	0.069	0.059	17	textual	=~	x8	3.359	-0.121	-0.120	-0.118
5	visual	=~	x5	7.441	-0.210	-0.189	-0.147	18	textual	=~	x9	4.796	0.138	0.137	0.136
6	visual	=~	x6	2.843	0.111	0.100	0.092	19	speed	=~	x1	0.014	0.024	0.015	0.013
7	visual	=~	x7	18.631	-0.422	-0.380	-0.349	20	speed	=~	x2	1.580	-0.198	-0.123	-0.105
8	visual	=~	x8	4.295	-0.210	-0.189	-0.187	21	speed	=~	x3	0.716	0.136	0.084	0.075
9	visual	=~	x9	36.411	0.577	0.519	0.515	22	speed	=~	x4	0.003	-0.005	-0.003	-0.003
10	textual	=~	x1	8.903	0.350	0.347	0.297	23	speed	=~	x5	0.201	-0.044	-0.027	-0.021
11	textual	=~	x2	0.017	-0.011	-0.011	-0.010	24	speed	=~	x6	0.273	0.044	0.027	0.025
12	textual	=~	x3	9.151	-0.272	-0.269	-0.238	25	speed	=~	x7	NA	NA	NA	NA
13	textual	=~	x4	NA	NA	NA	NA	26	speed	=~	x8	0.000	0.000	0.000	0.000
								27	speed	=~	x9	0.000	0.000	0.000	0.000

The fitted model

```
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

```
$mean
```

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

Examine the raw residuals

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

```
resid(fit)
```

```
$cov
```

	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	0.000								
x2	-0.041	0.000							
x3	-0.010	0.124	0.000						
x4	0.097	-0.017	-0.090	0.000					
x5	-0.014	-0.040	-0.219	0.008	0.000				
x6	0.077	0.038	-0.032	-0.012	0.005	0.000			
x7	-0.177	-0.242	-0.103	0.046	-0.050	-0.017	0.000		
x8	-0.046	-0.062	-0.013	-0.079	-0.047	-0.024	0.082	0.000	
x9	0.175	0.087	0.167	0.056	0.086	0.062	-0.042	-0.032	0.000

```
$mean
```

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

Examine the standardized residuals

```
fit <- cfa(HS.model, data = HolzingerSwineford1939)
resid(fit, type = "standardized")
```

\$cov

	x1	x2	x3	x4	x5	x6	x7	x8	x9
x1	0.000								
x2	-2.196	0.000							
x3	-1.199	2.692	0.000						
x4	2.465	-0.283	-1.948	NA					
x5	-0.362	-0.610	-4.443	0.856	NA				
x6	2.032	0.661	-0.701	NA	0.633	0.000			
x7	-3.787	-3.800	-1.882	0.839	-0.837	-0.321	NA		
x8	-1.456	-1.137	-0.305	-2.049	-1.100	-0.635	3.804	0.000	
x9	4.062	1.517	3.328	1.237	1.723	1.436	-2.771	NA	0.000

Many more examples

- The MPlus manual has data sets that may be explored with lavaan code.
 - Chapter 3: Regression and Path Analysis
 - Chapter 5: Confirmatory factor analysis and structural equation modeling
 - Chapter 6: Growth modeling
- LISREL manual also has suitable examples

SEM-AMOS wiki a warning

- It may seem odd to begin with a warning, but the popular misuse and misinterpretation of Structural Equation Modeling is so widespread that users of this wiki should be aware of some of the issues involved before they begin. While this warning is overly brief, you can follow-up these issues and more in the Further Reading section of this article.
- A number of these issues also apply to Confirmatory Factor Analysis. While Structural Equation Modeling has been popular in recent years to test the degree of fit between a proposed structural model and the emergent structure of the data, the perceived superiority of the technique is waning.
- Aside from the fact that the results of Structural Equation Modeling are often poorly reported, the conclusions drawn do not typically grasp the limitations of the technique.

SEM-AMOS wiki a warning page 2

- The most obvious, and some ways the most critical issue is that of incorrectly inferring a particular configuration of causal relationships from correlational data. This mistake can be illustrated with the simplest of all structural examples – that of 2 variables (variable A and B). If we ignore the additional complexity of latent structure, the number of possible causal structures is 4. Clearly, the number of possible models grows exponentially as the number of variables grows. In this example, the 4 possible causal models in this example are:
 - A causes B;
 - B causes A;
 - A and B cause each other (a recursive model);
 - finally, A and B are unrelated.

SEM-AMOS wiki warning page 3

- If A and B are indeed significantly correlated, it is likely that the first 3 models will be supported by significant fit statistics. If this is the case, what has been proven?
- Which of the 3 supported models is the correct model? What makes matters worse is that we have not even conclusively ruled out the last model. It is still possible that the correlation between A and B was spurious.
- To reinforce a maxim that most people know, but fail to apply to Structural Equation Modeling – you can not determine causation from correlation.
- Yet in most cases, researchers only test one or two models out of all the myriad of potential models, poorly report their results, then proclaim confirmation of their model (implying the exclusion of all other possible models).

SEM-AMOS wiki warning page 4

- So what is the value of Structural Equation Modeling?
- If large correlational datasets are already available, and a large range of plausible models are assessed, the results can be valuable in conceiving an experimental study that can test the proposed causal relationships.

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