

# Being negative can be good for your test

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4th Constrained Statistical Inference Meeting  
March 4, 2016 – Tilburg University

## lavaan 0.3-1 (first public version, May 2010)

Model converged normally after 35 iterations using ML

Minimum Function Chi-square	85.306
Degrees of freedom	24
P-value	0.0000

	Estimate	Std.err	Z-value	P(> z )
Latent variables:				
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
Latent covariances:				
visual ~~				
textual	0.408	0.074	5.552	0.000
speed	0.262	0.056	4.660	0.000
textual ~~				

<b>speed</b>	<b>0.173</b>	<b>0.049</b>	<b>3.518</b>	<b>0.000</b>
<b>Latent variances:</b>				
<b>visual</b>	<b>0.809</b>	<b>0.145</b>	<b>5.564</b>	<b>0.000</b>
<b>textual</b>	<b>0.979</b>	<b>0.112</b>	<b>8.737</b>	<b>0.000</b>
<b>speed</b>	<b>0.384</b>	<b>0.086</b>	<b>4.451</b>	<b>0.000</b>
<b>Residual variances:</b>				
<b>x1</b>	<b>0.549</b>	<b>0.114</b>	<b>4.833</b>	<b>0.000</b>
<b>x2</b>	<b>1.134</b>	<b>0.102</b>	<b>11.146</b>	<b>0.000</b>
<b>x3</b>	<b>0.844</b>	<b>0.091</b>	<b>9.317</b>	<b>0.000</b>
<b>x4</b>	<b>0.371</b>	<b>0.048</b>	<b>7.778</b>	<b>0.000</b>
<b>x5</b>	<b>0.446</b>	<b>0.058</b>	<b>7.642</b>	<b>0.000</b>
<b>x6</b>	<b>0.356</b>	<b>0.043</b>	<b>8.277</b>	<b>0.000</b>
<b>x7</b>	<b>0.799</b>	<b>0.081</b>	<b>9.823</b>	<b>0.000</b>
<b>x8</b>	<b>0.488</b>	<b>0.074</b>	<b>6.573</b>	<b>0.000</b>
<b>x9</b>	<b>0.566</b>	<b>0.071</b>	<b>8.003</b>	<b>0.000</b>

## lavaan 0.4-10 (Oct 2011)

Lavaan (0.4-10) converged normally after 41 iterations

Number of observations	301
Estimator	ML
Minimum Function Chi-square	85.306
Degrees of freedom	24
P-value	0.000

Parameter estimates:

Information				Expected
Standard Errors				Standard
	Estimate	Std.err	Z-value	P(> z )
Latent variables:				
visual =~				
x1	1.000			
x2	0.553	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:

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

## Variances:

x1	0.549	0.114
x2	1.134	0.102
x3	0.844	0.091
x4	0.371	0.048
x5	0.446	0.058
x6	0.356	0.043
x7	0.799	0.081
x8	0.488	0.074
x9	0.566	0.071
visual	0.809	0.145
textual	0.979	0.112
speed	0.384	0.086

## lavaan 0.5-19 (Oct 2015)

lavaan (0.5-19) converged normally after 35 iterations

Number of observations	301
Estimator	ML
Minimum Function Test Statistic	85.306
Degrees of freedom	24
P-value (Chi-square)	0.000

Parameter Estimates:

Information	Expected
Standard Errors	Standard

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

## 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

## asymptotic null distribution of likelihood-based tests

- simple hypothesis:  $H_0 : \theta = \theta_0$  versus  $H_a : \theta \neq \theta_0$
- Wald test, Score test, LR test with test statistic  $T_W, T_S, T_{LR}$
- it can be shown that ‘if regularity conditions hold’ all three test statistics follow a chi-square distribution under  $H_0$  if  $n \rightarrow \infty$
- for the LRT, the proof dates back to Wilks 1938
- what are these ‘regularity conditions’:
  - they are needed to establish the asymptotic normality of ML estimates
  - the one that matters here is:
    - the true parameter  $\theta$  (scalar or vector) must be an interior point of the parameter space  $\Theta$
  - in other words:  $\theta$  should NOT be a boundary point
- see for example Boos & Stefanski (2013): page 284 (scalar), page 286 (vector), page 291 (discussion)



## two nonstandard settings

- the alternative hypothesis is order-restricted
  - for example:

$$H_0 : \mu_1 = \mu_2 = \mu_3 \quad \text{versus} \quad H_a : \mu_1 < \mu_2 < \mu_3$$

- we need different test statistics; see Silvapulle and Sen (2005)
- the null hypothesis value,  $\theta_0$ , lies on the boundary of the parameter space, say  $\theta_0 = b$ 
  - the parameter space is restricted a priori (eg.  $\theta \geq b$ )
  - we violate the boundary assumption:  $\theta_0 = b$  is not an interior point
  - original reference: Chernoff (1954)
  - one-sided tests are a logical choice:

$$H_0 : \theta = \theta_0 \quad \text{versus} \quad H_a : \theta > \theta_0$$

## a typical example: inference on variance components

- consider a random-intercepts model:

$$Y_{ij} = x_{ij}^T \beta + b_i + \epsilon_{ij}$$

where  $Y_{ij}$  is the response for member  $j = 1, \dots, n_i$  of cluster  $i = 1, \dots, N$ ;  $x_{ij}$  is a vector of known covariate values;  $\beta$  is a vector of unknown regression coefficients; and  $b_i \sim N(0, \tau^2)$  is a cluster-specific random effect, independently distributed from the residual-error components  $\epsilon_{ij} \sim N(0, \sigma^2)$

- two-sided test:

$$H_0 : \tau^2 = 0 \quad \text{versus} \quad H_a : \tau^2 \neq 0$$

- if we assume the parameter space is restricted ( $\tau^2 \geq 0$ ), then 0 is on the boundary, and we can not use the classical test statistics

- one-sided test:

$$H_0 : \tau^2 = 0 \quad \text{versus} \quad H_a : \tau^2 > 0$$

- we can use special one-sided test statistics; Wald, Score and LR type statistics have been available for a while now

## interlude

- suppose you are fitting a random-intercept model (or any mixed model) using your favorite (but non-Bayesian) software
- you are interested in inference about variance components
- do you get a proper one-sided test?

## question

- what if ... variance components could be negative?

## negative variance components

- some of you may reject this immediately on mathematical (or religious, or philosophical) grounds
- but what if we treat those variance components in a more pragmatic way: as model parameters that are just part of a bigger story
- references in the world of SEM:
  - Kolenikov and Bollen (2012). Testing negative error variances: is a Heywood case a symptom of misspecification? *Sociological Methods & Research*, 41, 124–167
  - Savalei and Kolenikov (2008). Constrained versus unconstrained estimation in structural equation modeling. *Psychological Methods*, 13, 150–170.
- in the world of mixed models, negative variance (components) have been discussed by Nelder (1954), Thompson (1962), Searle, Casella and McCulloch (1992), Verbeke and Molenberghs (2000)

## mixed models: hierarchical versus marginal view

- the hierarchical model:

$$Y_{ij} = x_{ij}^T \beta + b_i + \epsilon_{ij}, \quad \text{where } b_i \sim N(0, \tau^2) \quad \text{and} \quad \epsilon_{ij} \sim N(0, \sigma^2)$$

- variances must be positive

- the marginal model:

- collecting all  $Y_{ij}$  from the same cluster into a (long) vector  $Y_i$
- the marginal distribution of  $Y_i$  is given by

$$Y_i \sim N(X_i \beta, V_i) \quad \text{where} \quad V_i = \tau^2 J_{n_i} + \sigma^2 I_{n_i}$$

- negative values for  $\tau^2$  (or  $\sigma^2$ ) are allowed, as long as  $V_i$  is positive definite
- but in this case, a hierarchical interpretation is no longer possible as there is no random-effects structure that would yield such a marginal model

## but there is more

- the hierarchical model can naturally be estimated using Bayesian estimation
- the hierarchical model can not be estimated in a frequentist framework: the random effects are treated as unobserved (latent) variables, and they must be integrated out
- so although we may ‘specify’ the model, as if it has a hierarchical structure, we always transform it to a marginal model
- consequently, at least from an estimation point of view, we can allow for negative variance components
- if we opt for this unconstrained, marginal approach
  - we may get negative variances
  - but standard tests (Wald, Score, LR) are no problem
  - negative variances can be informative (misspecification? negative intra-class correlation?)

## software aspects

- even if you adopt the marginal view, your software package may decide otherwise
- many popular software packages for fitting mixed models enforce strictly positive variance components (by default)
- if you are lucky, you can change the default, and ask for an unconstrained solution
- most SEM packages use unconstrained estimation per default
- as a user, you should be aware of the default options



## software defaults (variances)

software	constrained	unconstrained	user-controllable
HLM 6	x		
R lme	x		
R lmer	x		
MLwin	x (>level 1)	x (level 1)	x
SAS proc mixed	x		x
SPSS mixed	x		
Stata (xt)mixed	x		
AMOS		x	
EQS	x		x
LISREL		x	x
lavaan		x	x
Mplus		x	x
Stata sem	x		

## empirical example

- Alzheimer data (Hand & Taylor, 1987), but modified to get a negative variance for the random intercept
- 100 patients, 5 repeated measures of a score (number or words that a patient could correctly recall from a list); time is coded as 0,1,2,3,4
- model: linear growth curve with random intercept and random slope
- we can treat this as 2-level data (repeated measures nested within patients)
- to mimic MLM software, we constrained the residual variances of the 5 measures to be the same over time when using SEM software

## lavaan syntax

```
library(lavaan); Data <- read.csv("heywood_wide.csv")

model <- '
  i =~ 1*t1 + 1*t2 + 1*t3 + 1*t4 + 1*t5
  s =~ 0*t1 + 1*t2 + 2*t3 + 3*t4 + 4*t5

  t1 + t2 + t3 + t4 + t5 ~ 0*1

  # residual variance
  t1~~a*t1
  t2~~a*t2
  t3~~a*t3
  t4~~a*t4
  t5~~a*t5

  # fixed effects
  i ~ 1
  s ~ 1

  # variance components
  s ~~ s
  i ~~ i
  s ~~ i
'

fit <- sem(model, data = Data)
summary(fit)
```

## lavaan output

lavaan (0.5-19) converged normally after 42 iterations

Number of observations	100
Estimator	ML
Minimum Function Test Statistic	10.127
Degrees of freedom	14
P-value (Chi-square)	0.753

Parameter Estimates:

Information	Expected
Standard Errors	Standard

Latent Variables:

	Estimate	Std.Err	Z-value	P(> z )
i = ~				
t1	1.000			
t2	1.000			
t3	1.000			
t4	1.000			
t5	1.000			
s = ~				
t1	0.000			
t2	1.000			
t3	2.000			

t4	3.000
t5	4.000

## Covariances:

	Estimate	Std.Err	Z-value	P(> z )
i ~ s	0.329	0.183	1.796	0.073

## Intercepts:

	Estimate	Std.Err	Z-value	P(> z )
t1	0.000			
t2	0.000			
t3	0.000			
t4	0.000			
t5	0.000			
i	9.761	0.154	63.208	0.000
s	3.076	0.093	33.000	0.000

## Variances:

		Estimate	Std.Err	Z-value	P(> z )
t1	(a)	5.297	0.432	12.247	0.000
t2	(a)	5.297	0.432	12.247	0.000
t3	(a)	5.297	0.432	12.247	0.000
t4	(a)	5.297	0.432	12.247	0.000
t5	(a)	5.297	0.432	12.247	0.000
s		0.339	0.130	2.603	0.009
i		-0.794	0.425	-1.865	0.062

## Imer input

```
> library(lme4); Data <- read.table("heywood.dat", header = TRUE)
> fit <- lmer(y ~ 1 + time + (1 + time | id), data = Data, REML = FALSE)
> summary(fit)
```

```
Linear mixed model fit by maximum likelihood ['lmerMod']
Formula: y ~ 1 + time + (1 + time | id)
Data: Data
```

AIC	BIC	logLik	deviance	df.resid
2368.3	2393.6	-1178.2	2356.3	494

Scaled residuals:

Min	1Q	Median	3Q	Max
-2.64050	-0.64722	-0.03457	0.61837	3.01296

Random effects:

Groups	Name	Variance	Std.Dev.	Corr
id	(Intercept)	0.007527	0.08676	
	time	0.442050	0.66487	1.00
Residual		4.962683	2.22771	

Number of obs: 500, groups: id, 100

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	9.76050	0.17278	56.49
time	3.07582	0.09687	31.75

## overview

	$\beta_{00}$	$\beta_{10}$	$\sigma_{\epsilon}^2$	$\tau_0^2$	$\tau_1^2$	$\tau_{01}^2$
HLM	9.761 (.173)	3.076 (.098)	4.973 (np)	0.015 (np)	.448 (np)	0.057
lme	9.761 (.173)	3.076 (.098)	4.975 (np)	0.008 (np)	.459 (np)	0.043
lmer	9.761 (.173)	3.076 (.097)	4.975 (np)	0.008 (np)	.448 (np)	0.058
MLwin	9.761 (.173)	3.076 (.099)	4.966 (.351)	0.000 (-)	.481 (.092)	0.000
SAS	9.761 (.173)	3.076 (.097)	4.979 (.353)	0.000 (-)	.434 (.134)	0.081
SPSS	9.761 (.173)	3.076 (.096)	4.971 (.352)	0.000 (-)	.433 (.127)	0.083
Stata	9.761 (.173)	3.076 (.097)	4.963 (.353)	0.008 (.005)	.442 (.088)	0.058
AMOS	9.761 (.155)	3.076 (.094)	5.297 (.435)	-0.794 (.428)	0.339 (.131)	0.329
EQS	9.761 (.174)	3.076 (.097)	5.016 (.412)	0.000 (.494)	0.432 (.139)	0.081
lavaan	9.761 (.154)	3.076 (.093)	5.297 (.432)	-0.794 (.425)	0.339 (.130)	0.329
Mplus	9.761 (.154)	3.076 (.093)	5.297 (.432)	-0.794 (.425)	0.339 (.130)	0.329
LISREL	9.761 (.156)	3.076 (.094)	5.350 (.439)	-0.801 (.432)	0.343 (.132)	0.332
Stata	9.761 (.173)	3.076 (.098)	4.963 (.348)	0.001 (.000)	0.465 (.010)	0.026

## discussion

- is this old news?
- at least Molenberghs & Verbeke have discussed this in several papers in the (bio)statistics literature
- in the SEM world, we have a few papers (Savalei & Kolenikov, Kolenikov & Bollen)
- the discussion of ‘inference for variance components’ in handbooks written for applied users in psychology are confusing at best (if not plain wrong)
- should we (read: Leonard) write a paper about this?