

Example 4: CFA of Forgiveness of Situations ($N = 1103$) in Mplus v. 8.8, R lavaan 0.6-12, and a little bit of SAS MIXED (complete syntax and output available electronically)

This example comes from the Heartland Forgiveness Scale ([Yamhure Thompson et al., 2005](#)). Here we focus on the Forgiveness of Situations Subscale includes six items, three of which are reverse-coded, each on a 7-point scale:

1. When things go wrong for reasons that can't be controlled, I get stuck in negative thoughts about it. (R)
2. With time I can be understanding of bad circumstances in my life.
3. If I am disappointed by uncontrollable circumstances in my life, I continue to think negatively about them. (R)
4. I eventually make peace with bad situations in my life.
5. It's really hard for me to accept negative situations that aren't anybody's fault. (R)
6. Eventually I let go of negative thoughts about bad circumstances that are beyond anyone's control.

Response Anchors: 1 = Almost Always False of Me, 2=? , 3 = More Often False of Me, 4 = ?,

5 = More Often True of Me, 6 = ?, 7 = Almost Always True of Me

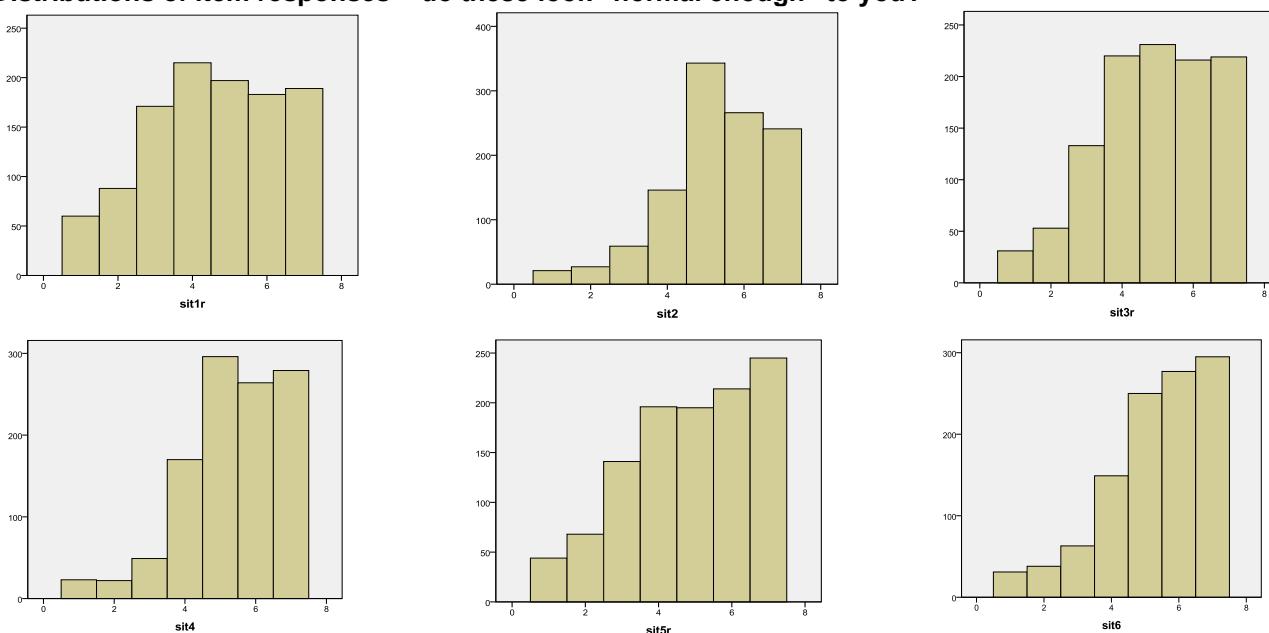
Observed Correlation Matrix	R1	2	R3	4	R5	6
R1	1.000					
2	0.240	1.000				
R3	0.647	0.317	1.000			
4	0.300	0.570	0.369	1.000		
R5	0.453	0.255	0.482	0.289	1.000	
6	0.297	0.457	0.356	0.448	0.304	1.000
Means	4.547	5.289	4.896	5.359	4.860	5.321
Variances	3.049	1.903	2.543	1.967	2.945	2.341

Observed Covariance Matrix	R1	2	R3	4	R5	6
R1	3.049					
2	0.577	1.903				
R3	1.802	0.697	2.543			
4	0.734	1.103	0.824	1.967		
R5	1.358	0.604	1.319	0.695	2.945	
6	0.795	0.965	0.868	0.962	0.798	2.341

To do CFA modeling, you only really need means, variances, and either correlations or covariances among items:

$$\text{Cov}(y_1, y_2) = \text{Cor}(y_1, y_2) * \text{SD}(y_1) * \text{SD}(y_2) \quad \text{OR} \quad \text{Cor}(y_1, y_2) = \text{Cov}(y_1, y_2) / \text{SD}(y_1) * \text{SD}(y_2)$$

Distributions of item responses – do these look “normal enough” to you?



Mplus Code to Read in Data and Select Options Across Models*(Note: DO NOT copy syntax from this handout—start from the original .inp file instead):*

```

TITLE:      CFA of Situation Factor
DATA:      FILE = Example4.csv;           ! Don't need path if in same directory
           FORMAT = free;             ! Default
           TYPE = INDIVIDUAL;        ! Default

VARIABLE:   NAMES = PersonID Self1 Self2r Self3 Self4r Self5 Self6r
           Other1r Other2r Other3r Other4r Other5r Other6r
           Sit1r Sit2r Sit3r Sit4r Sit5r Sit6r
           Selfsub Othsub Sitsub HFSsum;          ! Every variable in DATASET

           USEVARIABLES = Sit1r Sit2r Sit3r Sit4r Sit5r Sit6r;    ! Every variable in MODEL
           MISSING = ALL (99999);           ! Identify missing values
           IDVARIABLE = PersonID;         ! Identify person ID variable

ANALYSIS:   TYPE = GENERAL;           ! Default
           ESTIMATOR = MLR;            ! Robust ML

OUTPUT:    MODINDICES (6.635); ! Cheat codes to improve the model at p <.01 for df=1
           STDYX;                  ! Fully standardized solution
           RESIDUAL;                ! Standardized and normalized residuals for local fit

!SAVEDATA:  SAVE = FScores; FILE = FactorScores.dat; ! To save factor scores (optional)

!PLOT:      TYPE = PLOT1 PLOT2 PLOT3; ! To get plots (factor score distributions) as needed

MODEL:     ! (model syntax goes here, to be changed for each model as shown below)

```

Mplus Syntax for Model 1: Single Factor Using Fully Z-Scored Factor Scaling*(Factor Variance = 1, Factor Mean = 0, So All Item Loadings and Intercepts Estimated)***The following code refers to EVERY model parameter for completeness:**

```

! Model 1: Single Factor Using Fully Z-Scored Factor Scaling

! Item factor loadings --> @=fixed, *=free → * REQUIRED for first item if free
Sit BY Sit1r* Sit2r* Sit3r* Sit4r* Sit5r* Sit6r;

! Item intercepts --> [ ] indicates means or intercepts, @=fixed, *=free
[Sit1r* Sit2r* Sit3r* Sit4r* Sit5r* Sit6r];

! Item error variances --> just list item by itself, @=fixed, *=free
Sit1r* Sit2r* Sit3r* Sit4r* Sit5r* Sit6r;

! Factor variance fixed to 1 via @1 --> just list factor by itself, @=fixed, *=free
Sit@1;

! Factor mean fixed to 0 via @0 --> [ ] for means or intercepts, @=fixed, *=free
[Sit@0];

```

In reality, all you'd need to write to define this model is this:

```

! Item factor loadings --> @=fixed, *=free → * REQUIRED for first item if free
Sit BY Sit1r* Sit2r Sit3r Sit4r Sit5r Sit6r;

! Factor variance --> just list factor by itself, @=fixed, *=free
Sit@1;

```

By default, all intercepts are estimated separately and the factor mean is fixed at 0.

By default, all residual variances for the items are estimated separately, too.

By default, factor variances and covariances are estimated freely.

**Mplus Output for Model 1: Single Factor Using Fully Z-Scored Factor Scaling
(Factor Variance = 1, Factor Mean = 0, So All Item Loadings and Intercepts Estimated)**

UNSTANDARDIZED MODEL RESULTS

				Two-Tailed
	Estimate	S.E.	Est./S.E.	P-Value
FACTOR LOADINGS (regression slopes of item response on factor)				
SIT BY				
SIT1R	1.234	0.069	17.906	0.000
SIT2	0.702	0.074	9.441	0.000
SIT3R	1.241	0.063	19.846	0.000
SIT4	0.784	0.069	11.334	0.000
SIT5R	1.023	0.053	19.179	0.000
SIT6	0.819	0.069	11.942	0.000

Means (of Factor)

999 = "cannot be computed" - you will see 999 for any parameter that is FIXED

SIT	0.000	0.000	999.000	999.000
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Intercepts (of Items) - HERE, ARE ACTUAL ITEM MEANS BECAUSE FACTOR MEAN IS ZERO

SIT1R	4.547	0.053	86.474	0.000
SIT2	5.289	0.042	127.347	0.000
SIT3R	4.896	0.048	101.959	0.000
SIT4	5.359	0.042	126.895	0.000
SIT5R	4.860	0.052	94.060	0.000
SIT6	5.321	0.046	115.493	0.000

Variances (of Factor)

999 = "cannot be computed" - here, because the parameter is fixed to 1 already

SIT	1.000	0.000	999.000	999.000
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Residual Variances (variance of e's)

SIT1R	1.526	0.149	10.217	0.000
SIT2	1.409	0.128	11.014	0.000
SIT3R	1.004	0.135	7.456	0.000
SIT4	1.352	0.127	10.672	0.000
SIT5R	1.899	0.118	16.025	0.000
SIT6	1.671	0.159	10.517	0.000

Making use of the unstandardized model estimates:

Writing out the model—individual predicted values:

$$y_{1s} = \mu_1 + \lambda_1 F_s + e_{1s}$$

$$y_{1s} = 4.547 + 1.234 F_s + e_{1s}$$

Stay tuned! 1.523 will become the factor variance when item 1 is used as the "marker" item (whose loading is fixed to 1): 1.523 is the amount of item 1's variance that is due to the factor (with 1.526 error due to "not the factor" instead).

Writing out the model—predicted item variances and covariances:

$$Var(y_1) = (\lambda_1^2) Var(F) + Var(e_1)$$

$$Var(y_1) = (1.234^2)(1) + 1.526 = 1.523 + 1.526 = 3.049 (= \text{original item variance})$$

$$Cov(y_1, y_2) = \lambda_1 * Var(F) * \lambda_2$$

$$Cov(y_1, y_2) = 1.234 * 1 * 0.702 = 0.866$$

(actual covariance = 0.577, so the model over-predicted how related items 1 and 2 should be)

MPLUS STDYX STANDARDIZED MODEL RESULTS (FULLY STANDARDIZED WITH RESPECT TO X & Y)

			Two-Tailed	
	Estimate	S.E.	Est./S.E.	P-Value
FACTOR LOADINGS (correlations of item response with factor)				
Square these to get reliability (proportion "true variance") per item				
SIT	BY			
SIT1R	0.707	0.035	19.983	0.000
SIT2	0.509	0.053	9.545	0.000
SIT3R	0.778	0.034	22.655	0.000
SIT4	0.559	0.048	11.641	0.000
SIT5R	0.596	0.029	20.528	0.000
SIT6	0.535	0.047	11.392	0.000
Means (of Factor)				
SIT	0.000	0.000	999.000	999.000
Intercepts (of Items) → is intercept / SD(Y) → not usually reported				
SIT1R	2.604	0.057	45.888	0.000
SIT2	3.834	0.111	34.394	0.000
SIT3R	3.070	0.072	42.921	0.000
SIT4	3.821	0.111	34.441	0.000
SIT5R	2.832	0.066	43.095	0.000
SIT6	3.477	0.101	34.573	0.000
Variances (of Factor) → will always be 1 in a standardized solution				
SIT	1.000	0.000	999.000	999.000
Residual Variances (standardized variance of e residuals)				
SIT1R	0.500	0.050	10.009	0.000
SIT2	0.741	0.054	13.628	0.000
SIT3R	0.395	0.053	7.388	0.000
SIT4	0.687	0.054	12.786	0.000
SIT5R	0.645	0.035	18.619	0.000
SIT6	0.714	0.050	14.187	0.000
R-SQUARE (equals 1-residual variance OR standardized loading squared)				
SIT1R	0.500	0.050	9.991	0.000
SIT2	0.259	0.054	4.772	0.000
SIT3R	0.605	0.053	11.327	0.000
SIT4	0.313	0.054	5.821	0.000
SIT5R	0.355	0.035	10.264	0.000
SIT6	0.286	0.050	5.696	0.000

The standardized solution will look identical across methods of model identification with respect to the factor loadings, error variances, and R-square values for the items. The standardized intercepts will change because they depend on the unstandardized intercepts (but the standardized intercepts are rarely reported or interpreted anyway).

Making use of the standardized model estimates:

Writing out the model – predicted item correlations:

$$\text{Cor}(y_1, y_2) = \lambda_1 * \text{Var}(F) * \lambda_2$$

$$\text{Cor}(y_1, y_2) = .707 * 1 * .509 = .360$$

(actual correlation = .240, so the model over-predicted how related items 1 and 2 should be)

R Syntax for Model 1: Single Factor Using Fully Z-Scored Factor Scaling (Factor Variance = 1, Factor Mean = 0, So All Item Loadings and Intercepts Estimated)

The following code refers to EVERY model parameter for completeness:

```
# R Syntax for lavaan function: longer but more transparent version of model
Syntax1 =
# Define factor and request item factor loadings --> factor =~ item + item + item
  Sit =~ Sit1r + Sit2 + Sit3r + Sit4 + Sit5r + Sit6
# Item intercepts --> ~ 1 indicates means or intercepts
  Sit1r ~ 1; Sit2 ~ 1; Sit3r ~ 1; Sit4 ~ 1; Sit5r ~ 1; Sit6 ~ 1
# Item error variances or covariances --> use the ~~ command
  Sit1r ~~ Sit1r; Sit2 ~~ Sit2; Sit3r ~~ Sit3r; Sit4 ~~ Sit4; Sit5r ~~ Sit5r; Sit6 ~~ Sit6
# Factor variance fixed=1 and factor mean fixed=0 (* means fixed in lavaan)
  Sit ~~ 1*Sit; Sit ~ 0
#
# Use MLR estimation like in Mplus, z-scored latent variables (mean=0, SD=1)
  Model1 = lavaan(model=Syntax1, data=Example4, estimator="MLR", mimic="mplus", std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution
  summary(object=Model1, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)
```

In reality, all you'd need to write to define this model is this:

```
# R Syntax for sem or cfa functions: shorter but less transparent version of model
Syntax1short = "Sit =~ Sit1r + Sit2 + Sit3r + Sit4 + Sit5r + Sit6"
# Use MLR estimation like in Mplus, z-score latent factor (mean=0, SD=1)
  Model1short = sem(model=Syntax1short, data=Example4, estimator="MLR", mimic="mplus",
                     std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution (will be same as from lavaan)
  summary(object=Model1short, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)
```

R Lavaan Output for Model 1: Single Factor Using Fully Z-Scored Factor Scaling (Factor Variance = 1, Factor Mean = 0, So All Item Loadings and Intercepts Estimated)

Latent Variables:						
	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
Sit =~						
Sit1r	1.234	0.069	17.906	0.000	1.234	0.707
Sit2	0.702	0.074	9.441	0.000	0.702	0.509
Sit3r	1.241	0.063	19.847	0.000	1.241	0.778
Sit4	0.784	0.069	11.333	0.000	0.784	0.559
Sit5r	1.023	0.053	19.179	0.000	1.023	0.596
Sit6	0.819	0.069	11.942	0.000	0.819	0.535

For fully
standardized
results, use
std.all = STDYX

Intercepts:						
	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit1r	4.547	0.053	86.474	0.000	4.547	2.604
.Sit2	5.289	0.042	127.346	0.000	5.289	3.834
.Sit3r	4.896	0.048	101.959	0.000	4.896	3.070
.Sit4	5.359	0.042	126.896	0.000	5.359	3.821
.Sit5r	4.860	0.052	94.060	0.000	4.860	2.832
.Sit6	5.321	0.046	115.492	0.000	5.321	3.477
Sit	0.000				0.000	0.000

Variances:						
	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit1r	1.526	0.149	10.217	0.000	1.526	0.500
.Sit2	1.409	0.128	11.014	0.000	1.409	0.741
.Sit3r	1.004	0.135	7.456	0.000	1.004	0.395
.Sit4	1.352	0.127	10.672	0.000	1.352	0.687
.Sit5r	1.899	0.118	16.025	0.000	1.899	0.645
.Sit6	1.671	0.159	10.517	0.000	1.671	0.714
Sit	1.000				1.000	1.000

Next up: two equivalent ways of getting the same model, but with different scaling
(i.e., illustrating the results of different methods of identification, even though model fit is the same)

**Model 2. Single Factor Using Marker Item Loading = 1 and Factor Mean = 0
(Factor variance estimated given marker item loading=1, all intercepts estimated)**

```

! Mplus Model 2: Single Factor Scaled Using Marker Item Loading=1 and Factor Mean=0
! Factor variance estimated given marker item loading=1, all intercepts estimated
  Sit BY Sit1r@1 Sit2* Sit3r* Sit4* Sit5r* Sit6*;           ! Loadings (1st fixed=1)
  [Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*];                 ! Intercepts (all free)
  Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*;                   ! Residual variances (all free)
  Sit*;                                                       ! Factor variance (free)
  [Sit@0];                                                    ! Factor mean (fixed=0)

# R Syntax for lavaan function: longer but more transparent version of model
Syntax2 = "
# Define factor and request item factor loadings with first loading fixed to 1
  Sit =~ 1*Sit1r + Sit2 + Sit3r + Sit4 + Sit5r + Sit6
# Item intercepts all estimated
  Sit1r ~ 1; Sit2 ~ 1; Sit3r ~ 1; Sit4 ~ 1; Sit5r ~ 1; Sit6 ~ 1
# Item error variances all estimated
  Sit1r ~~ Sit1r; Sit2 ~~ Sit2; Sit3r ~~ Sit3r; Sit4 ~~ Sit4; Sit5r ~~ Sit5r; Sit6 ~~ Sit6
# Factor variance=estimated and factor mean=0
  " Sit ~~ Sit; Sit ~ 0

# Use MLR estimation like in Mplus, do not z-score latent factor (because variance is estimated)
  Model2 = lavaan(model=Syntax2, data=Example4, estimator="MLR", mimic="mplus", std.lv=FALSE)
# Print solution: get fit, get effect size, STDYX solution
  summary(object=Model2, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)

# R Syntax for sem or cfa functions: shorter but less transparent version of model
  Syntax2short = "Sit =~ 1*Sit1r + Sit2 + Sit3r + Sit4 + Sit5r + Sit6"
  Model2short = sem(model=Syntax2short, data=Example4, estimator="MLR", mimic="mplus",
                     std.lv=FALSE)

```

MPLUS UNSTANDARDIZED MODEL RESULTS

	Two-Tailed		
Estimate	S.E.	Est./S.E.	P-Value

FACTOR LOADINGS (regression slopes of item response on factor)

Here, loading for SIT1R is not tested because it is fixed=1

SIT	BY	Estimate	S.E.	Est./S.E.	P-Value
SIT1R		1.000	0.000	999.000	999.000
SIT2		0.569	0.083	6.830	0.000
SIT3R		1.005	0.035	28.555	0.000
SIT4		0.636	0.082	7.741	0.000
SIT5R		0.829	0.053	15.698	0.000
SIT6		0.664	0.081	8.143	0.000

In Mplus, fixed parameters have SE=0 and test statistics = 999 (so 999 is an undefined value)

Means (of Factor)

SIT	0.000	0.000	999.000	999.000
-----	-------	-------	---------	---------

Intercepts (of Items) - EXPECTED Y WHEN FACTOR = 0, or for mean of factor in sample

SIT1R	4.547	0.053	86.474	0.000
SIT2	5.289	0.042	127.347	0.000
SIT3R	4.896	0.048	101.960	0.000
SIT4	5.359	0.042	126.896	0.000
SIT5R	4.860	0.052	94.060	0.000
SIT6	5.321	0.046	115.492	0.000

Variances (of Factor)

SIT	1.523	0.170	8.954	0.000
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Residual Variances (variances of e's)

SIT1R	1.526	0.149	10.217	0.000
SIT2	1.409	0.128	11.014	0.000
SIT3R	1.004	0.135	7.456	0.000
SIT4	1.352	0.127	10.673	0.000
SIT5R	1.899	0.118	16.026	0.000
SIT6	1.671	0.159	10.517	0.000

R UNSTANDARDIZED MODEL RESULTS (TRUNCATED)

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
Sit =~						
Sit1r	1.000				1.234	0.707
Sit2	0.569	0.083	6.831	0.000	0.702	0.509
Sit3r	1.005	0.035	28.553	0.000	1.241	0.778
Sit4	0.636	0.082	7.741	0.000	0.784	0.559
Sit5r	0.829	0.053	15.698	0.000	1.023	0.596
Sit6	0.664	0.081	8.143	0.000	0.819	0.535

In R, fixed parameters have blanks for their SE and test statistics

Yet another equivalent alternative method for scaling the factor...**Model 3. Marker Item Loading = 1 and Intercept = 0 (Factor Variance and Mean Estimated)**

```

! Mplus Model 3: Single Factor Scaled Using Marker Item Loading=1 and Intercept=0
! Factor variance and mean estimated given marker item loading=1 and intercept=0
  Sit BY Sit1r@1 Sit2* Sit3r* Sit4* Sit5r* Sit6*;      ! Loadings (1st fixed=1)
  [Sit1r@0 Sit2* Sit3r* Sit4* Sit5r* Sit6*];          ! Intercepts (1st fixed=0)
  Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*;             ! Residual variances (all free)
  Sit*;                                                 ! Factor variance (free)
  [Sit*];                                              ! Factor mean (free)

# R Syntax for lavaan function: longer but more transparent version of model
Syntax3 =
# Define factor and request item factor loadings with first loading fixed to 1
  Sit =~ 1*Sit1r + Sit2 + Sit3r + Sit4 + Sit5r + Sit6
# Item intercepts: first is fixed to 0 and rest estimated
  Sit1r ~ 0; Sit2 ~ 1; Sit3r ~ 1; Sit4 ~ 1; Sit5r ~ 1; Sit6 ~ 1
# Item error variances all estimated
  Sit1r ~~ Sit1r; Sit2 ~~ Sit2; Sit3r ~~ Sit3r; Sit4 ~~ Sit4; Sit5r ~~ Sit5r; Sit6 ~~ Sit6
# Factor variance=estimated and factor mean=estimated
  " Sit ~ Sit; Sit ~ 1
# Use MLR estimation like in Mplus, do not z-score latent factor (mean and variance estimated)
  Model3 = lavaan(model=Syntax3, data=Example4, estimator="MLR", mimic="mplus", std.lv=FALSE)
# Print solution: get fit, get effect size, STDYX solution
  summary(object=Model3, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)

# No short version possible using sem that I know of

```

MPLUS UNSTANDARDIZED MODEL RESULTS (TRUNCATED)

Means (of Factor) → Note is mean of marker item 1
 SIT 4.547 0.053 86.474 0.000

Intercepts (of Items) - EXPECTED Y WHEN FACTOR = 0

HERE, WHICH IS WHEN ITEM 1 = 0 → beyond scale of item, so values are very low

SIT1R	0.000	0.000	999.000	999.000
SIT2	2.701	0.383	7.046	0.000
SIT3R	0.325	0.171	1.899	0.058
SIT4	2.469	0.380	6.504	0.000
SIT5R	1.092	0.246	4.431	0.000
SIT6	2.304	0.369	6.250	0.000

R UNSTANDARDIZED MODEL RESULTS (TRUNCATED)

Intercepts:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit1r	0.000				0.000	0.000
.Sit2	2.701	0.383	7.045	0.000	2.701	1.958
.Sit3r	0.325	0.171	1.899	0.058	0.325	0.204
.Sit4	2.469	0.380	6.502	0.000	2.469	1.760
.Sit5r	1.092	0.246	4.431	0.000	1.092	0.636
.Sit6	2.304	0.369	6.249	0.000	2.304	1.506
Sit	4.547	0.053	86.474	0.000	3.684	3.684

Calculating model degrees of freedom:

$$\text{Total DF} = \frac{v(v+1)}{2} + v = \frac{6(6+1)}{2} + 6 = 27$$

Spent: DF = 18

Leftover: Model DF = 9

Mplus model fit information for a single-factor model (same regardless of factor scaling method):

Number of Free Parameters 18 → is # of estimated parameters ("free" to be not 0)

Loglikelihood - use for testing differences in model fit across nested models

H0 Value	-11536.404	→ LL for YOUR specified model (bigger is better)
H0 Scaling Correction Factor for MLR	1.4159	→ indicates how far off from 1=normal
H1 Value	-11322.435	→ LL for a saturated (perfect) model
H1 Scaling Correction Factor for MLR	1.4073	→ indicates how far off from 1=normal

Information Criteria → "smaller is better" - use for nested or non-nested model comparisons

Akaike (AIC)	23108.808	→ AIC = (-2*LL _{H0}) + (2*estimated parameters)
Bayesian (BIC)	23198.912	→ BIC = (-2*LL _{H0}) + (LN N*estimated parameters)
Sample-Size Adjusted BIC (n* = (n + 2) / 24)	23141.739	→ BIC replacing N with (N + 2) / 24

Chi-Square Test of Model Fit (Significance is bad) → rescaled LRT for your model vs saturated

Value	307.799	
Degrees of Freedom	9	→ # parameters leftover = Model DF
P-Value	0.0000	
Scaling Correction Factor for MLR	1.3903	→ indicates how far off from normal=1 > 1 = leptokurtic distribution (too-fat tails) < 1 = platykurtotic distribution (too-thin tails)

- * The chi-square value for MLM, MLMV, **MLR**, ULSMV, WLSM and WLSMV cannot be used for chi-square difference testing in the regular way. MLM, MLR and WLSM chi-square difference testing is described on the Mplus website. MLMV, WLSMV, and ULSMV difference testing is done using the DIFFTEST option.

Where does this χ^2 value for "model fit" come from? A rescaled $-2\Delta LL$ model comparison of this one-factor model (H0) against the saturated model (H1) that perfectly reproduces the data covariances:

Step 1: Original $-2\Delta LL = -2(LL_{\text{fewer}} - LL_{\text{more}}) = -2(-11,536.404 + 11,322.435) = 427.938$

Step 2: Scaling correction = $[(\# \text{parms}_{\text{fewer}} * \text{scale}_{\text{fewer}}) - (\# \text{parms}_{\text{more}} * \text{scale}_{\text{more}})] / (\# \text{parms}_{\text{fewer}} - \# \text{parms}_{\text{more}})$
 $= [(18 * 1.4159) - (27 * 1.4073)] / (18 - 27) = -12.501 / -9 = 1.3903$

Step 3: Rescaled $-2\Delta LL = -2\Delta LL / \text{scaling correction} = 427.938 / 1.903 = 307.847 \rightarrow \sim \text{matches model } \chi^2$

Step 4: Difference in df = $\# \text{parms}_{\text{more}} - \# \text{parms}_{\text{fewer}} = 27 - 18 = 9$

RMSEA (Root Mean Square Error Of Approximation) → how much worse than 0 = saturated model

Estimate	0.173	→ smaller is better
90 Percent C.I.	0.157	0.190 → CI will become smaller as sample size increases
Probability RMSEA <= .05	0.000	→ so RMSEA does NOT overlap .05 (is signif > .05)

CFI/TLI → how much better than 0 = null model (bigger is better)

CFI	0.732
TLI	0.553

SRMR (Standardized Root Mean Square Residual) → how much worse than 0 = saturated model

Value	0.086
-------	-------

Chi-Square Test of Model Fit for the Baseline Model → LRT of null vs saturated (who cares)

Value	1128.693
Degrees of Freedom	15
P-Value	0.0000

Where does this χ^2 value for “fit of the baseline model” come from? A rescaled -2LL model comparison of the independence “null” model with NO covariances to the saturated model:

Step 1: Original $-2\Delta LL = -2*(LL_{\text{fewer}} - LL_{\text{more}}) = -2(-12,312.952 + 11,322.435) = 1,981.034$

Step 2: Scaling correction = [(#parms_{fewer}*scale_{fewer}) - (#parms_{more}*scale_{more})] / (#parms_{fewer} - #parms_{more})
 $= [(12 * 0.9725) - (27 * 1.4073)] / (12 - 27) = -26.372 / -15 = 1.7551$

Step 3: Rescaled $-2\Delta LL = -2\Delta LL / \text{scaling correction} = 1,981.034 / 1.7551 = 1,128.704 \rightarrow \sim \text{matches baseline } \chi^2$

Step 4: Difference in df = #parms_{more} - #parms_{fewer} = 27 - 12 = 15

What's the point? This baseline model fit test tells us whether there are any covariances at all (i.e., whether it even makes sense to try to fit latent factors to predict the item covariances).

R model fit information for a single-factor model (same regardless of factor scaling method):

Model Test User Model:

	For ML:	For MLR:
Test Statistic	427.937	307.803 → LRT for your model vs H1
Degrees of freedom	9	9 → # parms leftover = Model DF
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.390 → how far off from normal=1

Yuan-Bentler correction (Mplus variant)

Model Test Baseline Model:

Test statistic	1981.034	1128.693 → LRT of null vs saturated
Degrees of freedom	15	15
P-value	0.000	0.000
Scaling correction factor		1.755

User Model versus Baseline Model: → how much better than 0 = null model (bigger is better)

Comparative Fit Index (CFI)	0.787	0.732
Tucker-Lewis Index (TLI)	0.645	0.553

Robust Comparative Fit Index (CFI)	0.787	→ not in Mplus
Robust Tucker-Lewis Index (TLI)	0.646	→ not in Mplus

Loglikelihood and Information Criteria: → For LL, bigger is better; for IC, smaller is better

Loglikelihood user model (H0)	-11536.404	-11536.404 → LL for your model
Scaling correction factor		1.416 → how far off from 1=normal

for the MLR correction		
Loglikelihood unrestricted model (H1)	-11322.435	-11322.435 → LL for saturated model
Scaling correction factor		1.407 → how far off from 1=normal
for the MLR correction		

Akaike (AIC)	23108.808	23108.808 → smaller is better
Bayesian (BIC)	23198.912	23198.912
Sample-size adjusted Bayesian (BIC)	23141.739	23141.739

Root Mean Square Error of Approximation: → how much worse than 0 = saturated model

RMSEA	0.205	0.173 → smaller is better
90 Percent confidence interval - lower	0.189	0.160
90 Percent confidence interval - upper	0.222	0.188
P-value RMSEA <= 0.05	0.000	0.000

Robust RMSEA	0.205 → not in Mplus
90 Percent confidence interval - lower	0.185
90 Percent confidence interval - upper	0.224

Standardized Root Mean Square Residual: → how much worse than 0 = saturated model

SRMR	0.086	0.086
------	-------	-------

FYI for demonstration purposes, here is how to fit the saturated (Unstructured) Baseline Model: Item means, variances, and covariances in original data

```

! Saturated Model for Demonstration Purposes
! All item means, variances, and covariances estimated
  ! Item means --> [ ] indicates means or intercepts, @=fixed, *=free
    [Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*];
  ! Item variances --> just list item by itself, @=fixed, *=free
    Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*;
  ! Item covariances --> just list all by all, @=fixed, *=free
    Sit1r  Sit2  Sit3r  Sit4  Sit5r  Sit6  WITH
      Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*;

```

```

# R Syntax for lavaan function: longer but more transparent version of model
SyntaxSat =
# Item means all estimated
Sit1r ~ 1; Sit2 ~ 1; Sit3r ~ 1; Sit4 ~ 1; Sit5r ~ 1; Sit6 ~ 1
# Item variances all estimated
Sit1r ~~ Sit1r; Sit2 ~~ Sit2; Sit3r ~~ Sit3r; Sit4 ~~ Sit4; Sit5r ~~ Sit5r; Sit6 ~~ Sit6
# Covariances all estimated
Sit1r ~~ Sit2; Sit1r ~~ Sit3r; Sit1r ~~ Sit4; Sit1r ~~ Sit5r; Sit1r ~~ Sit6
Sit2 ~~ Sit3r; Sit2 ~~ Sit4; Sit2 ~~ Sit5r; Sit2 ~~ Sit6; Sit3r ~~ Sit4; Sit3r ~~ Sit5r
" Sit3r ~~ Sit6; Sit4 ~~ Sit5r; Sit4 ~~ Sit6; Sit5r ~~ Sit6

```

```

ModelSat = lavaan(model=SyntaxSat, data=Example4, estimator="MLR", mimic="mplus")
summary(ModelSat, fit.measures=TRUE, standardized=TRUE)
# Get saturated model-implied means, variances, and covariances
fitted(object=ModelSat)

```

Model fit information for the saturated model: illustrating what the χ^2 test of global model fit means

Number of Free Parameters 27 → all possible means, variances, covariances

Loglikelihood		Note that H0 and H1 are now the same! Our H0 model = the H1 saturated model.	
H0 Value	-11322.435		
H0 Scaling Correction Factor for MLR	1.4073		
H1 Value	-11322.435	From R: > fitted(object = ModelSat)	
H1 Scaling Correction Factor for MLR	1.4073	\$cov	
Information Criteria			
Akaike (AIC)	22698.870	Sit1r Sit2 Sit3r Sit4 Sit5r Sit6	
Bayesian (BIC)	22834.027	Sit1r 3.049	
Sample-Size Adjusted BIC (n* = (n + 2) / 24)	22748.268	Sit2 0.577 1.903	
Chi-Square Test of Model Fit		Sit3r 1.802 0.697 2.543	
Value	0.000*	Sit4 0.734 1.103 0.824 1.967	
Degrees of Freedom	0	Sit5r 1.358 0.604 1.319 0.695 2.945	
P-Value	0.0000	Sit6 0.795 0.965 0.868 0.962 0.798 2.341	
Scaling Correction Factor for MLR	1.0000	 \$mean	
		Sit1r Sit2 Sit3r Sit4 Sit5r Sit6	
		4.547 5.289 4.896 5.359 4.860 5.321	

As another FYI for demonstration purposes, here is how to fit the Independence (Null) Baseline Model: Item means and variances, but NO covariances

```

! Null Independence Model for Demonstration Purposes
! All means and variances estimated, all covariances fixed=0
  ! Item means --> [ ] indicates means or intercepts, @=fixed, *=free
    [Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*];
  ! Item variances --> just list item by itself, @=fixed, *=free
    Sit1r* Sit2* Sit3r* Sit4* Sit5r* Sit6*;
  ! NO Item covariances --> just list all by all, @=fixed to 0
    Sit1r  Sit2  Sit3r  Sit4  Sit5r  Sit6  WITH
      Sit1r@0 Sit2@0 Sit3r@0 Sit4@0 Sit5r@0 Sit6@0;

# R Syntax for lavaan function: longer but more transparent version of model
SyntaxNull =
# Item means all estimated
  Sit1r ~ 1; Sit2 ~ 1; Sit3r ~ 1; Sit4 ~ 1; Sit5r ~ 1; Sit6 ~ 1
# Item variances all estimated
  " Sit1r ~~ Sit1r; Sit2 ~~ Sit2; Sit3r ~~ Sit3r; Sit4 ~~ Sit4; sit5r ~~ sit5r; Sit6 ~~ Sit6

ModelNull = lavaan(model=SyntaxNull, data=Example4, estimator="MLR", mimic="mplus")
summary(ModelNull, fit.measures=TRUE)
# Get null model-implied means, variances, and covariances
fitted(object=ModelNull)

```

Model fit for the independence “null” model: illustrating what RMSEA, CFI, and TLI mean

Number of Free Parameters	12
Loglikelihood	
H0 Value	-12312.952
H0 Scaling Correction Factor for MLR	0.9725
H1 Value	-11322.435
H1 Scaling Correction Factor for MLR	1.4073
Information Criteria	
Akaike (AIC)	24649.904
Bayesian (BIC)	24709.974
Sample-Size Adjusted BIC (n* = (n + 2) / 24)	24671.859
Chi-Square Test of Model Fit	
Value	1128.692*
Degrees of Freedom	15
P-Value	0.0000
Scaling Correction Factor for MLR	1.7552
RMSEA (Root Mean Square Error Of Approximation)	
Estimate	0.259
90 Percent C.I.	0.247 0.272
Probability RMSEA <= .05	0.000
CFI/TLI	
CFI	0.000
TLI	0.000
Chi-Square Test of Model Fit for the Baseline Model	
Value	1128.693
Degrees of Freedom	15
P-Value	0.0000
SRMR (Standardized Root Mean Square Residual)	
Value	0.300

```

From R: > fitted(object = ModelNull)
$cov
  Sit1r Sit2 Sit3r Sit4 Sit5r Sit6
Sit1r 3.049
Sit2 0.000 1.903
Sit3r 0.000 0.000 2.543
Sit4 0.000 0.000 0.000 1.967
Sit5r 0.000 0.000 0.000 0.000 2.945
Sit6 0.000 0.000 0.000 0.000 0.000 2.341

$mean
  Sit1r Sit2 Sit3r Sit4 Sit5r Sit6
4.547 5.289 4.896 5.359 4.860 5.321

```

Note that the model fit is the same as the “baseline” model fit given before.

Although not 0, this is the worst possible RMSEA while still allowing separate means and variances per item in these data. RMSEA is a parsimony-corrected absolute fit index (so, its fit is relative to the saturated model).

CFI and TLI are 0 because they are “incremental fit” indices relative to the independence model (which this is).

SRMR is also an absolute fit index (relative to saturated model), so this is the worst it gets for these data, too.

Here is what the single-factor model implied for our item means, variances, and covariances:

Mplus: scroll to RESIDUAL OUTPUT → info in “Model Estimated” sections

R: # R Get model-implied means, variances, and covariances
fitted(object=Model1)

```
$cov → Model-implied matrix
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
Sit1r 3.049
Sit2  0.867 1.903
Sit3r 1.531 0.872 2.543
Sit4  0.968 0.551 0.973 1.967
Sit5r 1.263 0.719 1.269 0.803 2.945
Sit6  1.011 0.575 1.016 0.642 0.838 2.341
```

```
$mean → Model-implied means
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
Sit1r 4.547 5.289 4.896 5.359 4.860 5.321
```

VS Saturated Model fitted(object=ModelSat)

```
$cov → From original data
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
Sit1r 3.049
Sit2  0.577 1.903
Sit3r 1.802 0.697 2.543
Sit4  0.734 1.103 0.824 1.967
Sit5r 1.358 0.604 1.319 0.695 2.945
Sit6  0.795 0.965 0.868 0.962 0.798 2.341
```

```
$mean → From original data
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
Sit1r 4.547 5.289 4.896 5.359 4.860 5.321
```

But global fit for the one-factor model is not good enough—let’s examine the residual info, which provides the discrepancy between what the model implies and what the H1 data model says:

R Get residual (discrepancy) info for local misfit in covariance and correlation form
resid(object=Model1, type="raw"); resid(object=Model1, type="cor")

R \$cov → Discrepancies in means, variances, and covariances

```
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
Sit1r 0.000
Sit2 -0.290 0.000
Sit3r 0.271 -0.174 0.000
Sit4 -0.234 0.552 -0.149 0.000
Sit5r 0.096 -0.114 0.050 -0.108 0.000
Sit6 -0.216 0.390 -0.149 0.319 -0.040 0.000

$mean
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
  0     0     0     0     0     0
```

All values are from: observed – predicted

Top: The variances and means have discrepancies=0 → perfect recovery, which means they are not the cause of our bad fit. Thus, misfit results from the difference between the observed and model-implied (model-recreated) covariances.

R \$cov → Discrepancies in correlations (STDX)

```
  Sit1r Sit2  Sit3r Sit4  Sit5r Sit6
Sit1r 0.000
Sit2 -0.120 0.000
Sit3r 0.097 -0.079 0.000
Sit4 -0.096 0.285 -0.067 0.000
Sit5r 0.032 -0.048 0.018 -0.045 0.000
Sit6 -0.081 0.185 -0.061 0.149 -0.015 0.000
```

Bottom: Although this is called the “residuals for correlations” matrix in Mplus, it is NOT the same as “residual correlations”, which are error correlations as model parameters. These are **discrepancies** in correlation!

From Mplus: Residuals for Correlations (Observed – Predicted Correlations)

	SIT1R	SIT2	SIT3R	SIT4	SIT5R	SIT6
SIT1R	0.000					
SIT2	-0.120	0.000				
SIT3R	0.097	-0.079	0.000			
SIT4	-0.096	0.285	-0.067	0.000		
SIT5R	0.032	-0.048	0.018	-0.045	0.000	
SIT6	-0.081	0.185	-0.061	0.149	-0.015	0.000

We can also get “normalized” residuals, which can be thought of as z-scores for how large the residual leftover covariance is in absolute terms. Because the denominator decreases with sample size, however, these values may be inflated in large samples, so look for *relatively* large values.

R: Get normalized residuals (like z-scores for discrepancies)
resid(object=Model1, type="normalized")

“Normalized” Residuals for Inter-Item Covariances: $z = (\text{observed} - \text{predicted}) / \text{SE}(\text{observed})$

Mplus Normalized Residuals for Covariances (same results in R)					
	SIT1R	SIT2	SIT3R	SIT4	SIT5R
SIT1R	0.000				
SIT2	-3.503	0.000			
SIT3R	2.977	-2.253	0.000		
SIT4	-2.928	6.560	-1.959	0.000	
SIT5R	0.960	-1.434	0.548	-1.372	0.000
SIT6	-2.345	4.721	-1.756	3.925	-0.444
					0.000

NEGATIVE NORMALIZED RESIDUAL → Less related than you predicted (want to be less together)
POSITIVE NORMALIZED RESIDUAL → More related than you predicted (want to be more together)

Why might the normalized residuals (leftover correlations) for the positive-worded items be larger than for the negatively-worded items?

What do the “cheat code” modification indices suggest we do to improve model fit?

MPLUS: MODEL MODIFICATION INDICES (MI = change in model test chi-square)

Minimum M.I. value for printing the modification index 6.635

EPC = EXPECTED PARAMETER CHANGE

M.I. E.P.C. Std E.P.C. StdYX E.P.C.

WITH Statements (SUGGESTED ERROR COVARIANCES for unaccounted for multidimensionality)

SIT2	WITH SIT1R	49.618	-0.464	-0.464	-0.316
SIT3R	WITH SIT1R	143.624	1.023	1.023	0.827
SIT3R	WITH SIT2	34.877	-0.357	-0.357	-0.300
SIT4	WITH SIT1R	36.280	-0.403	-0.403	-0.280
SIT4	WITH SIT2	161.318	0.702	0.702	0.509
SIT4	WITH SIT3R	29.202	-0.336	-0.336	-0.288
SIT6	WITH SIT1R	24.079	-0.358	-0.358	-0.224
SIT6	WITH SIT2	63.893	0.486	0.486	0.317
SIT6	WITH SIT3R	22.386	-0.319	-0.319	-0.246
SIT6	WITH SIT4	46.541	0.415	0.415	0.276

Get “cheat code” modification indices
 modificationindices(object = Model14, sort.=TRUE)

```
> modificationindices(object = Model11, sort. = TRUE)
   lhs op   rhs      mi     epc    sepc.lv sepc.all sepc.nox
 27 Sit2 ~~ Sit4 224.295  0.702   0.702   0.509   0.509
 22 Sit1r ~~ Sit3r 199.679  1.023   1.023   0.827   0.827
 29 Sit2 ~~ Sit6  88.835  0.486   0.486   0.317   0.317
 21 Sit1r ~~ Sit2 68.985 -0.464  -0.464  -0.316  -0.316
 34 Sit4 ~~ Sit6 64.710  0.415   0.415   0.276   0.276
 23 Sit1r ~~ Sit4 50.442 -0.403  -0.403  -0.280  -0.280
 26 Sit2 ~~ Sit3r 48.505 -0.357  -0.357  -0.300  -0.300
 30 Sit3r ~~ Sit4 40.616 -0.336  -0.336  -0.288  -0.288
 25 Sit1r ~~ Sit6 33.475 -0.358  -0.358  -0.224  -0.224
 32 Sit3r ~~ Sit6 31.132 -0.319  -0.319  -0.246  -0.246
 28 Sit2 ~~ Sit5r 7.117 -0.151  -0.151  -0.092  -0.092
 33 Sit4 ~~ Sit5r 6.906 -0.149  -0.149  -0.093  -0.093
 24 Sit1r ~~ Sit5r 6.417  0.176   0.176   0.104   0.104
 31 Sit3r ~~ Sit5r 3.548  0.123   0.123   0.089   0.089
 35 Sit5r ~~ Sit6 0.736 -0.053  -0.053  -0.030  -0.030
```

The MI column provides the non-robust (regular ML) version of the change in model test chi-square from adding that parameter. Lavaan used to provide the robust version as mi.scaled, but not anymore...?

But we come to the same conclusion either way about which new parameters would result in the biggest change in fit... note that all suggestions are error covariances! This is not a coincidence, because they cannot suggest a new model!

These results suggest that wording valence is playing a larger role in the pattern of covariance across items than what the one-factor model predicts. Rather than adding many covariances among the residuals for specific items, how about a two-factor model based on wording instead?

Model 4. Model with Two Fully Z-Scored Factors (adding labels to the end of each parameter to use in computing omega reliability per factor)

```

! Mplus Model 4: Two Factors Using Fully Z-Scored Factor Scaling with all parameters labeled
  SitP BY Sit2*  Sit4*  Sit6*  (L1-L3);          ! SitP loadings (all free)
  SitN BY Sit1r* Sit3r* Sit5r* (L4-L6);          ! SitN loadings (all free)
  [Sit2*  Sit4*  Sit6*]  (I1-I3);                ! SitP intercepts (all free)
  [Sit1r* Sit3r* Sit5r*] (I4-I6);                ! SitN intercepts (all free)
  Sit2*  Sit4*  Sit6*  (E1-E3);                ! SitP residual variances (all free)
  Sit1r* Sit3r* Sit5r* (E4-E6);                ! SitN residual variances (all free)
  SitP@1 (VarP);  SitN@1 (VarN);                ! Factor variances (fixed=1)
  SitP WITH SitN* (FactCov);                  ! Factor covariance (free)
  [SitP@0 SitN@0] (MeanP MeanN);                ! Factor means (fixed=0)

MODEL CONSTRAINT: ! Mplus syntax to compute omega sum score reliability per factor
NEW(OmegaP OmegaN); ! Using 1 as placeholder for factor variances
OmegaP = (1*(L1+L2+L3)**2) / ((1*(L1+L2+L3)**2) + (E1+E2+E3));
OmegaN = (1*(L4+L5+L6)**2) / ((1*(L4+L5+L6)**2) + (E4+E5+E6));
! Don't forget to include any error covariances, too! (Lecture 4 slide 77)

# R Syntax for lavaan function: longer but more transparent version of model
Syntax4 = " ! Note labels are in order of inclusion, not according to item name
# Define factor and request item factor loadings all estimated
  SitP =~ L1*Sit2 + L2*Sit4 + L3*Sit6      # Pos items
  SitN =~ L4*Sit1r + L5*Sit3r + L6*Sit5r    # Neg items
# Item intercepts all estimated
  Sit2 ~ I1*1; Sit4 ~ I2*1; Sit6 ~ I3*1    # Pos items
  Sit1r ~ I4*1; Sit3r ~ I5*1; Sit5r ~ I6*1  # Neg items
# Item error variances all estimated
  Sit2 ~~ E1*Sit2; Sit4 ~~ E2*Sit4; Sit6 ~~ E3*Sit6    # Pos items
  Sit1r ~~ E4*Sit1r; Sit3r ~~ E5*Sit3r; Sit5r ~~ E6*Sit5r # Neg items
# Factor variances fixed=1 and factor means fixed=0 (won't allow labels)
  SitP ~~ 1*SitP; SitN ~~ 1*SitN; SitP ~ 0; SitN ~ 0
# Factor covariance estimated (and labeled)
  SitP ~~ FactCov*SitN
# Calculate Omega Reliability for Sum Score Per Factor (1=factor variance):
OmegaP := (1*(L1 + L2 + L3)^2) / ( (1*(L1 + L2 + L3)^2) + (E1 + E2 + E3))
OmegaN := (1*(L4 + L5 + L6)^2) / ( (1*(L4 + L5 + L6)^2) + (E4 + E5 + E6))
# Don't forget to include any error covariances, too! (Lecture 4 slide 77)

"
# Use MLR estimation like in Mplus, z-score latent factors (mean=0, SD=1)
Model4 = lavaan(model=Syntax4, data=Example4, estimator="MLR", mimic="mplus", std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution
summary(object=Model4, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)

print("LRT for one factor vs two factors: is correlation < 1?")
anova(Model1, Model4)

# Get model-implied means, variances, and covariances
fitted(object=Model4)
# Get residual (discrepancy) info for local misfit in covariance and correlation form
resid(object=Model4, type="raw"); resid(object=Model4, type="cor")
# Get normalized residuals (like z-scores for discrepancies)
resid(object=Model4, type="normalized")
# Get "cheat code" modification indices
modificationindices(object = Model4, sort.=TRUE)

# R Syntax for sem or cfa functions: shorter but less transparent version of model
Syntax4short =
  SitP =~ L1*Sit2 + L2*Sit4 + L3*Sit6      # Pos items
  SitN =~ L4*Sit1r + L5*Sit3r + L6*Sit5r    # Neg items
# Use MLR estimation like in Mplus, z-score latent factor (mean=0, SD=1)
Model4short = sem(model=Syntax4short, data=Example4, estimator="MLR", mimic="mplus",
                 std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution (will be same as from lavaan)
summary(object=Model4short, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)

```

MPLUS MODEL FIT INFORMATION

Number of Free Parameters	19
Loglikelihood	
H0 Value	-11340.140
H0 Scaling Correction Factor for MLR	1.4017
H1 Value	-11322.435
H1 Scaling Correction Factor for MLR	1.4073
Information Criteria	
Akaike (AIC)	22718.281
Bayesian (BIC)	22813.391
Sample-Size Adjusted BIC (n* = (n + 2) / 24)	22753.042
Chi-Square Test of Model Fit	
Value	24.924*
Degrees of Freedom	8
P-Value	0.0016
Scaling Correction Factor for MLR	1.4207
RMSEA (Root Mean Square Error Of Approximation)	
Estimate	0.044
90 Percent C.I.	0.025 0.064
Probability RMSEA <= .05	0.667
CFI/TLI	
CFI	0.985
TLI	0.972

Chi-Square Test of Model Fit for the Baseline Model	
Value	1128.693
Degrees of Freedom	15
P-Value	0.0000

SRMR (Standardized Root Mean Square Residual)	
Value	0.029

FILL IN				CALCULATED				
Models: FewerParms in Row 1 MoreParms in Row 2	Test of -2ΔLL Difference							
	Model H0 LL	H0 LL Scale Factor	# Free Parms	Diff in LL * -2	Diff Scaling Correction	Scaled Diff in -2LL	DF Diff	Exact P-Value
Demonstration of LRTs you have to do yourself (alternative H0 models)								
One-Factor	-11,536.404	1.4158	18					
Two-Factor	-11,340.140	1.4017	19					
Test of Difference				392.528	1.1479	341.953	1	0.0000

Model comparison from lavaan using anova:

```
[1] "LRT for one factor vs two factors: is correlation < 1?"
> anova(Model1, Model4)
Scaled Chi-Squared Difference Test (method = "satorra.bentler.2001")
lavaan NOTE:
```

The "Chisq" column contains standard test statistics, not the robust test that should be reported per model. A robust difference test is a function of two standard (not robust) statistics.

Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
Model4	8	22718	22813	35.4		
Model1	9	23109	23199	427.9	1	<2e-16

Is the 2-factor model better than the 1-factor model? How do we know?

Rescaled likelihood ratio test between H0 models (-2LL rescaled difference test):

1. $-2\Delta LL = -2^* \text{ difference in LL:}$

$$-2^*(-11,536.404 - -11,340.140) = 392.528$$

2. difference scaling correction:

$$\frac{(\text{parms}_1 * \text{scale}_1) - (\text{parms}_2 * \text{scale}_2)}{(\text{parms}_1 - \text{parms}_2)} / (18 * 1.4158) - (19 * 1.4017) / (18 - 19) = 1.1479$$

3. rescaled difference = $-2\Delta LL / \text{scaling correction:}$

$$392.528 / 1.1479 = 341.953$$

4. compare rescaled difference to χ^2 with DF = ADF:

critical χ^2 for DF = 1 is 3.84, so because 341.953 is > 3.84, the model fit significantly improved

Report LRT as: $-2\Delta LL(1) = 342, p < .001$
Or as: $\chi^2(1) = 342, p < .001$

MPLUS UNSTANDARDIZED RESULTS

		Estimate	S.E.	Two-Tailed	
SITP	BY			Est./S.E.	P-Value
SIT2		1.007	0.052	19.487	0.000
SIT4		1.064	0.050	21.195	0.000
SIT6		0.956	0.053	18.203	0.000
SITN	BY				
SIT1R		1.325	0.048	27.698	0.000
SIT3R		1.349	0.044	30.514	0.000
SIT5R		1.009	0.055	18.358	0.000

SITP WITH SITN = factor covariance (= correlation if variances=1)

	0.564	0.041	13.776	0.000
--	-------	-------	--------	-------

Means

SITP	0.000	0.000	999.000	999.000
SITN	0.000	0.000	999.000	999.000

Intercepts

SIT1R	4.547	0.053	86.474	0.000
SIT2	5.289	0.042	127.347	0.000
SIT3R	4.896	0.048	101.959	0.000
SIT4	5.359	0.042	126.896	0.000
SIT5R	4.860	0.052	94.060	0.000
SIT6	5.321	0.046	115.492	0.000

Variances

SITP	1.000	0.000	999.000	999.000
SITN	1.000	0.000	999.000	999.000

Residual Variances

SIT1R	1.294	0.103	12.547	0.000
SIT2	0.888	0.097	9.173	0.000
SIT3R	0.724	0.092	7.857	0.000
SIT4	0.835	0.093	9.003	0.000
SIT5R	1.926	0.119	16.128	0.000
SIT6	1.428	0.134	10.684	0.000

New/Additional Parameters

OMEGAP	0.744	0.020	37.956	0.000
OMEGAN	0.775	0.014	56.803	0.000

STDYX STANDARDIZED RESULTS

		Estimate	S.E.	Two-Tailed	
SITP	BY			Est./S.E.	P-Value
SIT2		0.730	0.032	22.794	0.000
SIT4		0.759	0.029	25.995	0.000
SIT6		0.625	0.035	17.949	0.000
SITN	BY				
SIT1R		0.759	0.022	34.072	0.000
SIT3R		0.846	0.021	39.657	0.000
SIT5R		0.588	0.030	19.651	0.000
SITP	WITH				
SITN		0.564	0.041	13.776	0.000
SITP	WITH				
SIT1R		0.425	0.034	12.567	0.000
SIT2		0.467	0.047	9.976	0.000
SIT3R		0.285	0.036	7.895	0.000
SIT4		0.425	0.044	9.589	0.000
SIT5R		0.654	0.035	18.576	0.000
SIT6		0.610	0.043	14.029	0.000
R-SQUARE					
SIT1R		0.575	0.034	17.036	0.000
SIT2		0.533	0.047	11.397	0.000
SIT3R		0.715	0.036	19.829	0.000
SIT4		0.575	0.044	12.998	0.000
SIT5R		0.346	0.035	9.826	0.000
SIT6		0.390	0.043	8.974	0.000

Omega =

$$\text{Var(Factor)} * (\text{Sum of loadings})^2 / \text{Var(Factor)} * (\text{Sum of loadings})^2 + \text{Sum of error variances} + 2 * \text{Sum of error covariances}$$

Omega for Positive Factor = .744

$$1.0 * (1.007 + 1.064 + 0.956)^2 / 1.0 * (1.007 + 1.064 + 0.956)^2 + (0.888 + 0.835 + 1.428) + 2 * 0$$

(alpha was .746, btw)

Omega for Negative Factor = .775

$$1.0 * (1.325 + 1.349 + 1.009)^2 / 1.0 * (1.325 + 1.349 + 1.009)^2 + (1.294 + 0.724 + 1.926) + 2 * 0$$

(alpha was .780, btw)

R Output, which adds the labels you gave to the parameters:**Latent Variables:**

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP =~							
Sit2	(L1)	1.007	0.052	19.487	0.000	1.007	0.730
Sit4	(L2)	1.064	0.050	21.195	0.000	1.064	0.759
Sit6	(L3)	0.956	0.053	18.203	0.000	0.956	0.625
SitN =~							
Sit1r	(L4)	1.325	0.048	27.698	0.000	1.325	0.759
Sit3r	(L5)	1.349	0.044	30.514	0.000	1.349	0.846
Sit5r	(L6)	1.009	0.055	18.358	0.000	1.009	0.588

Covariances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP ~~							
SitN	(FctC)	0.564	0.041	13.775	0.000	0.564	0.564

Intercepts:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2	(I1)	5.289	0.042	127.346	0.000	5.289	3.834
.Sit4	(I2)	5.359	0.042	126.896	0.000	5.359	3.821
.Sit6	(I3)	5.321	0.046	115.492	0.000	5.321	3.477
.Sit1r	(I4)	4.547	0.053	86.474	0.000	4.547	2.604
.Sit3r	(I5)	4.896	0.048	101.959	0.000	4.896	3.070
.Sit5r	(I6)	4.860	0.052	94.060	0.000	4.860	2.832
SitP		0.000				0.000	0.000
SitN		0.000				0.000	0.000

Variances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2	(E1)	0.888	0.097	9.173	0.000	0.888	0.467
.Sit4	(E2)	0.835	0.093	9.003	0.000	0.835	0.425
.Sit6	(E3)	1.428	0.134	10.684	0.000	1.428	0.610
.Sit1r	(E4)	1.294	0.103	12.547	0.000	1.294	0.425
.Sit3r	(E5)	0.724	0.092	7.857	0.000	0.724	0.285
.Sit5r	(E6)	1.926	0.119	16.128	0.000	1.926	0.654
SitP		1.000				1.000	1.000
SitN		1.000				1.000	1.000

R-Square:

	Estimate
Sit2	0.533
Sit4	0.575
Sit6	0.390
Sit1r	0.575
Sit3r	0.715
Sit5r	0.346

Defined Parameters:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
OmegaP	0.744	0.020	37.956	0.000	0.744	0.748
OmegaN	0.775	0.014	56.802	0.000	0.775	0.779

The two-factor model appears to fit well overall, but is there any remaining sizeable local misfit?

```
# R Get model-implied means, variances, and covariances
fitted(object=Model14)
# R Get residual (discrepancy) info for local misfit in covariance and correlation form
resid(object=Model14, type="raw"); resid(object=Model14, type="cor")
# R Get normalized residuals (like z-scores for discrepancies)
resid(object=Model14, type="normalized")
# R Get "cheat code" modification indices
modificationindices(object = Model14, sort.=TRUE)
```

Mplus residuals of correlation matrix (observed – model-implied correlation):

	SIT1R	SIT2	SIT3R	SIT4	SIT5R	SIT6
SIT1R	0.000					
SIT2	-0.073	0.000				
SIT3R	0.006	-0.031	0.000			
SIT4	-0.025	0.016	0.007	0.000		
SIT5R	0.007	0.013	-0.015	0.037	0.000	
SIT6	0.030	0.001	0.057	-0.026	0.097	0.000

Mplus “Normalized” residuals (z-like statistic for how far off each covariance is):

	SIT1R	SIT2	SIT3R	SIT4	SIT5R	SIT6
SIT1R	0.000					
SIT2	-2.125	0.000				
SIT3R	0.172	-0.896	0.000			
SIT4	-0.768	0.370	0.192	0.000		
SIT5R	0.212	0.382	-0.464	1.128	0.000	
SIT6	0.869	0.031	1.658	-0.676	2.847	0.000

Any suggested cheat codes? (only available in Mplus when not using MODEL CONSTRAINT)**MPLUS MODEL MODIFICATION INDICES**

Minimum M.I. value for printing the modification index 6.635

M.I. E.P.C. Std E.P.C. StdYX E.P.C.

BY Statements – these are cross-loadings

SITN	BY SIT2	9.775	-0.224	-0.224	-0.162
SITN	BY SIT6	10.828	0.245	0.245	0.160

WITH Statements – these are error covariances

SIT4	WITH SIT2	10.830	0.332	0.332	0.386
SIT6	WITH SIT4	9.773	-0.273	-0.273	-0.250

```
> modificationindices(object = Model4, sort. = TRUE)
   lhs op   rhs    mi    epc sepc.lv sepc.all sepc.nox
31 SitN =~ Sit6 15.383 0.245  0.245  0.160  0.160
32 Sit2 ~~ Sit4 15.383 0.332  0.332  0.386  0.386
37 Sit4 ~~ Sit6 13.886 -0.273 -0.273 -0.250 -0.250
29 SitN =~ Sit2 13.885 -0.224 -0.224 -0.162 -0.162
26 SitP =~ Sit1r 9.244 -0.223 -0.223 -0.128 -0.128
46 Sit3r ~~ Sit5r 9.244 -0.277 -0.277 -0.234 -0.234
```

=~ are cross-loadings
~~ are error covariances

Because we have no real theoretical or defendable reason to fit any of these suggested parameters, we will not add any new parameters. This will be about as good as it gets.

Let's examine the estimated distribution of the factor scores for each factor... these are EAP (expected a posteriori) estimates, which are the mean of each person's random factor distribution

To do so in Mplus: turn on SAVE and PLOT options in syntax; open *Plot* menu in output, select *View plots*, select *histograms*, select *view*, in *plot properties* tab use the drop-down menu to select your factor name, then go to *display properties* tab and select *histogram/density plot*. When the plot appears, you can customize the axes using the menus visible by right-clicking on the plot.

To do so in R: see example code online, with more details in this handout:

https://jonathantemplin.com/wp-content/uploads/2017/09/EPSY906_Example04.nb_.html#

MPLUS SAMPLE STATISTICS FOR ESTIMATED FACTOR SCORES**SAMPLE STATISTICS****Means**

	SITP	SITP_SE	SITN	SITN_SE
1	0.000	0.472	0.000	0.418

Covariances

	SITP	SITP_SE	SITN	SITN_SE
SITP	0.777			
SITP_SE	0.000	0.000		
SITN	0.533	0.000	0.825	
SITN_SE	0.000	0.000	0.000	0.000

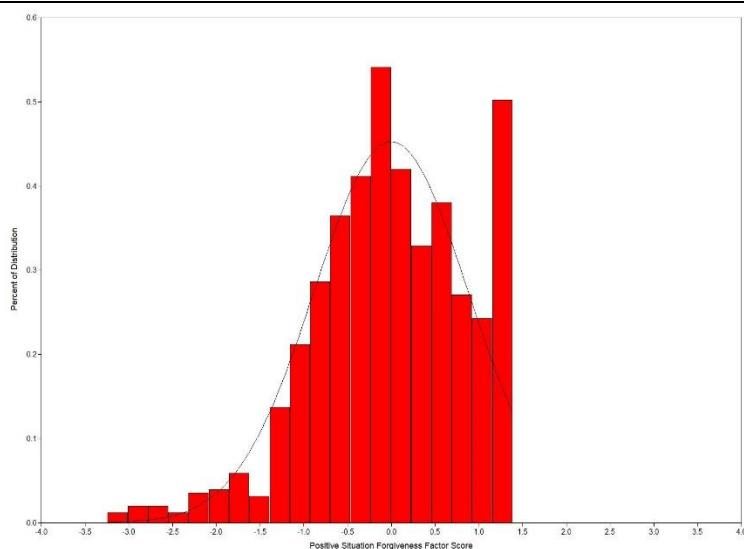
Correlations

	SITP	SITP_SE	SITN	SITN_SE
SITP	1.000			
SITP_SE	999.000	1.000		
SITN	0.665	999.000	1.000	
SITN_SE	999.000	999.000	999.000	1.000

Positive factor score SE = 0.472
Negative factor score SE = 0.418

Although the variance of each factor was supposed to be 1.0, the variance of the factor scores is < 1.0 because of shrinkage (positive factor var = 0.777; negative factor var = 0.825).

Likewise, the correlation between the factors was .56, but the correlation between the estimated factor scores is .67 instead (given the shrinkage).



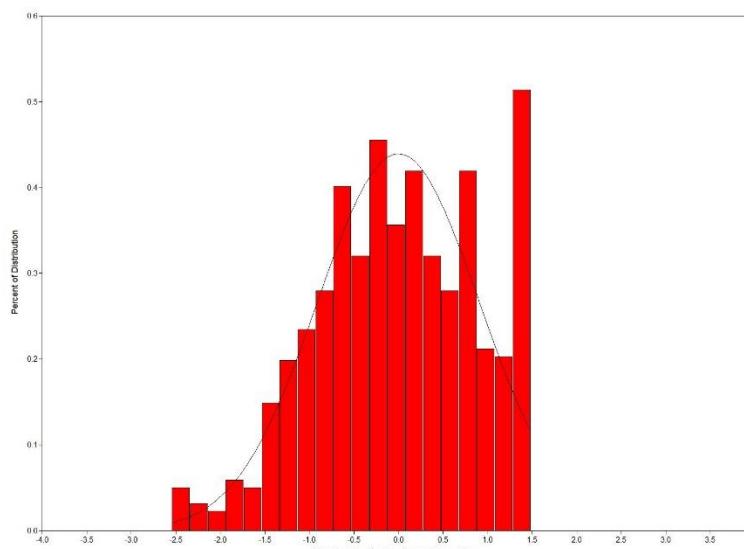
The **positive factor scores** have an estimated mean of 0 with a variance of 0.78 instead of the model-based 1.00.

The SE for each person's factor score is 0.472. Treating factor scores as observed variables is like saying SE = 0.

Positive factor score =
Score $\pm 2 \times 0.472$ = Score ± 0.944 !

Positive items factor score reliability =

$$\frac{\sigma_F^2}{\sigma_F^2 + SE_{FS}^2} = \frac{1}{1+0.472^2} = .818$$



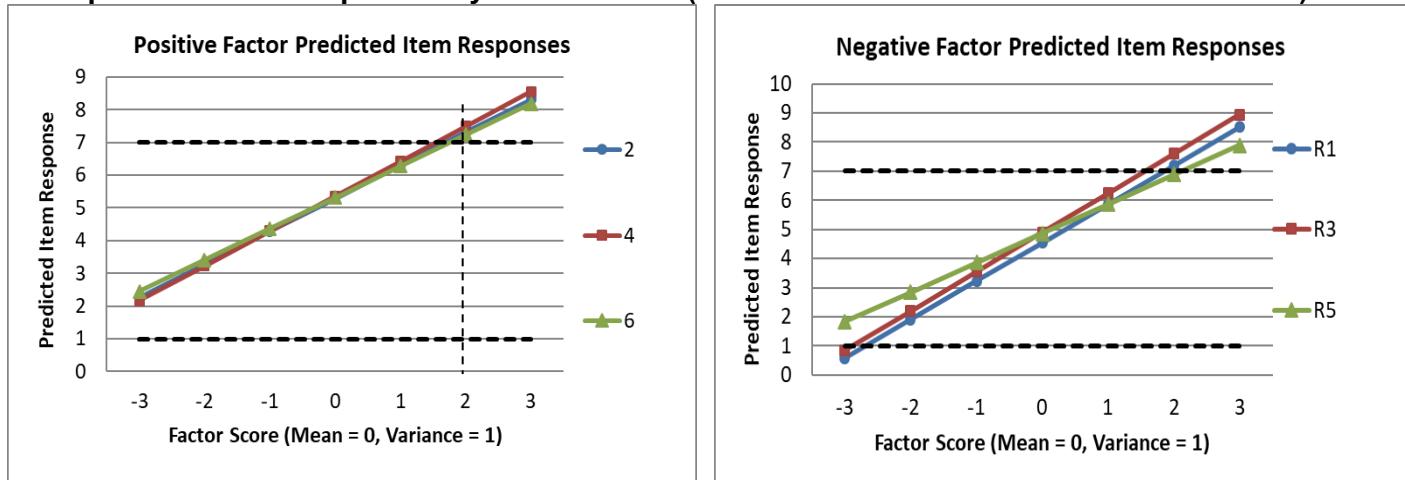
The negative factor scores have an estimated mean of 0 with a variance of 0.825 instead of the model-based 1.00.

The SE for each person's factor score is 0.418, so ± 0.836 !

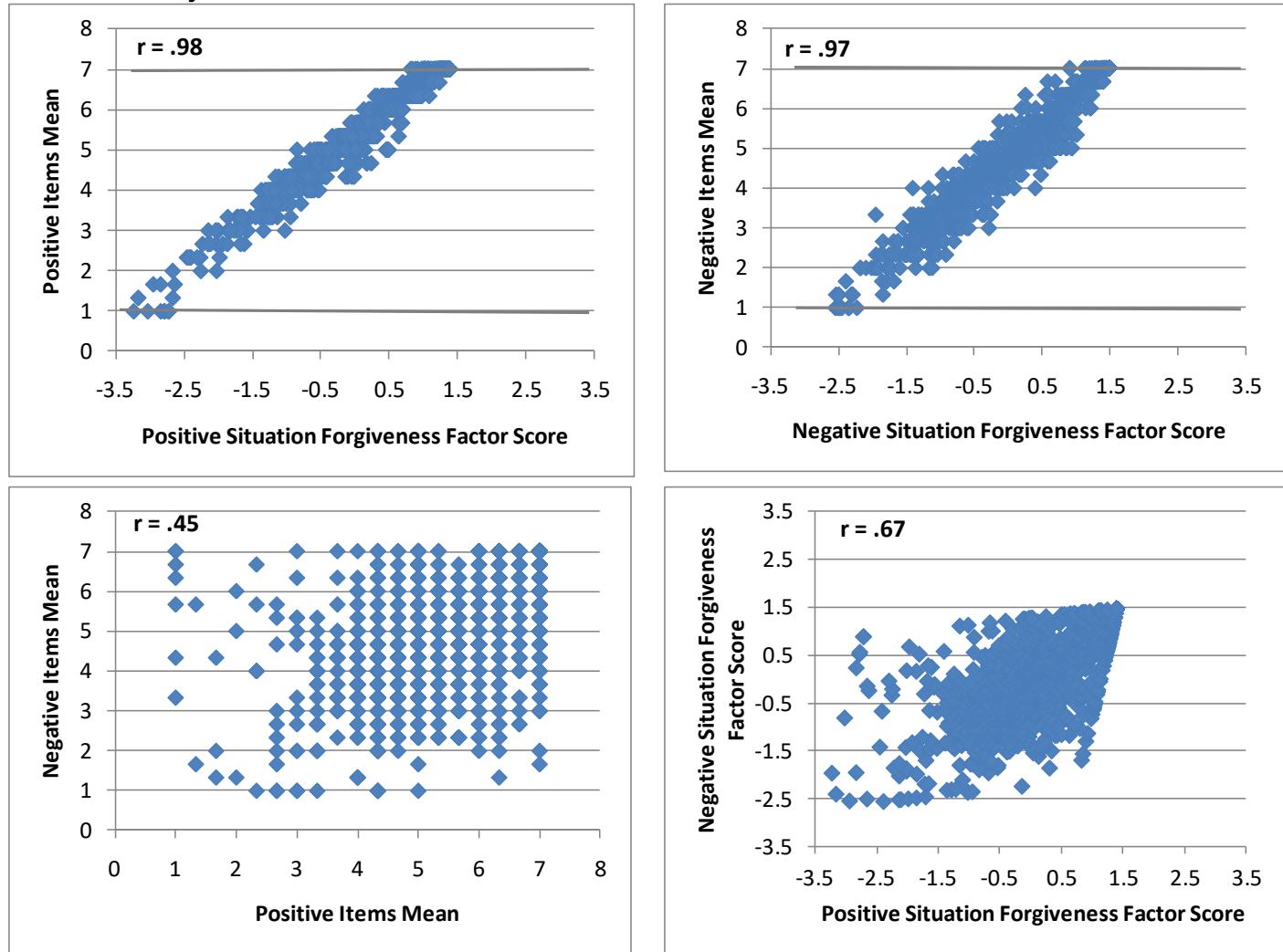
Negative items factor score reliability =

$$\frac{\sigma_F^2}{\sigma_F^2 + SE_{FS}^2} = \frac{1}{1+0.418^2} = .851$$

The negative factor scores retain more variance (and have a smaller SE) because there is more information in them, due to higher factor loadings (greater reliability) of their items.

Model-predicted item responses by factor scores (see excel sheet “Factor Model Predictions”):


What if we had just taken the mean of the three items for each subscale?



There are problems with either of these observed variable approaches: The **mean of the items** appears to have less variability (i.e., fewer possible scores), and any sum or mean across items assumes that all items should be weighted equally and have no error. The **estimated factor scores** do not have the same properties as estimated for the factor in the model (i.e., less variance for each factor, higher correlation among the factors). **So use SEM if possible!**

Another example: Formal Tests of CTT Assumptions

We will test the CTT assumption of tau-equivalence (equal factor loadings), one factor at a time.

If those hold, we can then test the assumption of parallel items (equal error variances, too).

First, in Model 5 we will test “tau-equivalence” (equal loadings) of the negative factor only:

```

! Mplus Model 5: Two Factors with Tau-Equivalent Negative Items Only
  SitP BY Sit2* Sit4 Sit6;                      ! SitP loadings (all free)
  SitN BY Sit1r* Sit3r Sit5r (NegLoad);        ! SitN loadings (all held equal)
  SitP@1; SitN@1;                               ! Factor variances (fixed=1)
  SitP WITH SitN*;                            ! Factor covariance (free)

# Model 5: Two Factors with Tau-Equivalent Negative Items Only (loadings equal)
Syntax5 = " ! Note labels are in order of inclusion, not according to item name
# Define factor and request item factor Loadings all estimated
  SitP =~ L1*Sit2 + L2*Sit4 + L3*Sit6          # Pos items
  SitN =~ NegLoad*Sit1r + NegLoad*Sit3r + NegLoad*Sit5r    # Neg items
"
# Use MLR estimation like in Mplus, z-score latent factors (mean=0, SD=1)
Model15 = sem(model=Syntax5, data=Example4, estimator="MLR", mimic="mplus", std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution
summary(object=Model15, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)

print("LRT: Does the shared loading for the negative items make fit worse?")
anova(Model15, Model14)

```

R lavaan output:

Latent Variables:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP =~							
Sit2	(L1)	1.007	0.052	19.491	0.000	1.007	0.730
Sit4	(L2)	1.063	0.050	21.201	0.000	1.063	0.758
Sit6	(L3)	0.957	0.052	18.258	0.000	0.957	0.626
SitN =~							
Sit1r	(NgLd)	1.254	0.032	38.957	0.000	1.254	0.735
Sit3r	(NgLd)	1.254	0.032	38.957	0.000	1.254	0.805
Sit5r	(NgLd)	1.254	0.032	38.957	0.000	1.254	0.682

Covariances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP ~~	SitN	0.575	0.041	13.856	0.000	0.575	0.575

Intercepts:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2		5.289	0.042	127.346	0.000	5.289	3.834
.Sit4		5.359	0.042	126.896	0.000	5.359	3.821
.Sit6		5.321	0.046	115.492	0.000	5.321	3.477
.Sit1r		4.547	0.053	86.474	0.000	4.547	2.667
.Sit3r		4.896	0.048	101.959	0.000	4.896	3.141
.Sit5r		4.860	0.052	94.060	0.000	4.860	2.644
SitP		0.000				0.000	0.000
SitN		0.000				0.000	0.000

Variances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2		0.889	0.096	9.216	0.000	0.889	0.467
.Sit4		0.837	0.092	9.046	0.000	0.837	0.425
.Sit6		1.425	0.134	10.630	0.000	1.425	0.609
.Sit1r		1.335	0.083	16.149	0.000	1.335	0.459
.Sit3r		0.857	0.069	12.337	0.000	0.857	0.353
.Sit5r		1.806	0.115	15.715	0.000	1.806	0.535
SitP		1.000				1.000	1.000
SitN		1.000				1.000	1.000

Why are the standardized factor loadings for the negative factor not held equal like the unstandardized loadings are?

Mplus Model Fit:

Fit of previous 2-factor model as baseline:		Fit of tau-equivalent negative items 2-factor model:	
Number of Free Parameters	19	Number of Free Parameters	17
Loglikelihood		Loglikelihood	
H0 Value	-11340.140	H0 Value	-11357.612
H0 Scaling Correction Factor for MLR	1.4017	H0 Scaling Correction Factor for MLR	1.4474
H1 Value	-11322.435	H1 Value	-11322.435
H1 Scaling Correction Factor for MLR	1.4073	H1 Scaling Correction Factor for MLR	1.4073
RMSEA (Root Mean Square Error Of Approximation)		RMSEA (Root Mean Square Error Of Approximation)	
Estimate	0.044	Estimate	0.062
90 Percent C.I.	0.025 0.064	90 Percent C.I.	0.046 0.079
Probability RMSEA <= .05	0.667	Probability RMSEA <= .05	0.102
CFI/TLI		CFI/TLI	
CFI	0.985	CFI	0.962
TLI	0.972	TLI	0.943

Does the assumption of tau-equivalence hold for the negative items? Let's see the lavaan anova output:

```
[1] "LRT: Does the shared loading for the negative items make fit worse?"
> anova(Model5, Model4)
Scaled Chi-Squared Difference Test (method = "satorra.bentler.2001")
lavaan NOTE:
  The "Chisq" column contains standard test statistics, not the
  robust test that should be reported per model. A robust difference
  test is a function of two standard (not robust) statistics.

      Df   AIC   BIC Chisq Chisq diff Df diff  Pr(>Chisq)
Model4  8 22718 22813  35.4
Model5 10 22749 22834  70.3      34.5        2 0.000000032
```

Second, Model 6 tests tau-equivalence of the factor loadings for the positive factor only:

```
! Mplus Model 6: Two Factors with Tau-Equivalent Positive Items Only
  SitP BY Sit2* Sit4 Sit6 (PosLoad);      ! SitP loadings (all held equal)
  SitN BY Sit1r* Sit3r Sit5r;            ! SitN loadings (all free)
  Sit2  Sit4  Sit6           (E1-E3);     ! SitP residual variances (all free)
  Sit1r Sit3r Sit5r;                   ! SitN residual variances (all free)
  SitP@1; SitN@1;                     ! Factor variances (fixed=1)
  SitP WITH SitN*;                  ! Factor covariance (free)

MODEL CONSTRAINT:
  NEW(AlphaP);    ! This is now equivalent to alpha reliability for pos items
  AlphaP = (1*(PosLoad*3)**2) / ((1*(PosLoad*3)**2) + (E1+E2+E3));

# Model 6: Two Factors with Tau-Equivalent Positive Items Only (loadings equal)
Syntax6 = " ! Note labels are in order of inclusion, not according to item name
# Define factor and request item factor loadings all estimated
  SitP =~ PosLoad*Sit2 + PosLoad*Sit4 + PosLoad*Sit6      # Pos items
  SitN =~ L4*Sit1r + L5*Sit3r + L6*Sit5r                 # Neg items
# Item error variances all estimated
  Sit2 ~~ E1*Sit2; Sit4 ~~ E2*Sit4; Sit6 ~~ E3*Sit6      # Pos items
  Sit1r ~~ E4*Sit1r; Sit3r ~~ E5*Sit3r; Sit5r ~~ E6*Sit5r # Neg items
# Calculate Alpha Reliability for Sum Score of Positive Factor:
  AlphaP := ((PosLoad*3)^2) / ( ((PosLoad*3)^2) + (E1 + E2 + E3))
#
# Use MLR estimation like in Mplus, z-score latent factors (mean=0, SD=1)
Model6 = sem(model=Syntax6, data=Example4, estimator="MLR", mimic="mplus", std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution
summary(object=Model6, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)
```

```
print("LRT: Does the shared loading for the positive items make fit worse?")
anova(Model16, Model14)
```

R lavaan output:

Latent Variables:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP ==							
Sit2	(PsLd)	1.014	0.036	28.388	0.000	1.014	0.734
Sit4	(PsLd)	1.014	0.036	28.388	0.000	1.014	0.733
Sit6	(PsLd)	1.014	0.036	28.388	0.000	1.014	0.653
SitN ==							
Sit1r	(L4)	1.325	0.048	27.726	0.000	1.325	0.759
Sit3r	(L5)	1.349	0.044	30.531	0.000	1.349	0.846
Sit5r	(L6)	1.010	0.055	18.369	0.000	1.010	0.588

Covariances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP ~~							
SitN		0.567	0.040	14.130	0.000	0.567	0.567

Intercepts:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2		5.289	0.042	127.346	0.000	5.289	3.828
.Sit4		5.359	0.042	126.896	0.000	5.359	3.873
.Sit6		5.321	0.046	115.492	0.000	5.321	3.426
.Sit1r		4.547	0.053	86.474	0.000	4.547	2.604
.Sit3r		4.896	0.048	101.959	0.000	4.896	3.070
.Sit5r		4.860	0.052	101.959	0.000	4.860	2.832
SitP		0.000				0.000	0.000
SitN		0.000				0.000	0.000

Variances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2	(E1)	0.881	0.083	10.587	0.000	0.881	0.461
.Sit4	(E2)	0.886	0.075	11.767	0.000	0.886	0.463
.Sit6	(E3)	1.384	0.118	11.737	0.000	1.384	0.574
.Sit1r	(E4)	1.295	0.103	12.580	0.000	1.295	0.425
.Sit3r	(E5)	0.725	0.092	7.873	0.000	0.725	0.285
.Sit5r	(E6)	1.925	0.119	16.117	0.000	1.925	0.654
SitP		1.000				1.000	1.000
SitN		1.000				1.000	1.000

Defined Parameters:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
AlphaP		0.746	0.020	38.200	0.000	0.746	0.764

```
[1] "LRT: Does the shared loading for the positive items make fit worse?"
```

```
> anova(Model16, Model14)
```

Scaled Chi-Squared Difference Test (method = "satorra.bentler.2001")

lavaan NOTE:

The "Chisq" column contains standard test statistics, not the robust test that should be reported per model. A robust difference test is a function of two standard (not robust) statistics.

Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
Model14	8	22718	22813	35.4		
Model16	10	22718	22803	38.7	2.6	0.27

Mplus model fit:

Number of Free Parameters	17
Loglikelihood	
H0 Value	-11341.773
H0 Scaling Correction Factor	1.4187
for MLR	
H1 Value	-11322.435
H1 Scaling Correction Factor	1.4073
for MLR	

Does the assumption of tau-equivalence hold for the positive items? How do we know?

Given that tau-equivalence held for the positive factor, in Model 7 we can also test the assumption of parallel items as equal residual variances (in addition to equal factor loadings):

```

! Mplus Model 7: Two Factors with Parallel Positive Items Only
SitP BY Sit2* Sit4 Sit6 (PosLoad); ! SitP loadings (all held equal)
SitN BY Sit1r* Sit3r Sit5r; ! SitN loadings (all free)
Sit2 Sit4 Sit6 (PosError); ! SitP residual variances (all held equal)
Sit1r Sit3r Sit5r; ! SitN residual variances (all free)
SitP@1; SitN@1; ! Factor variances (fixed=1)
SitP WITH SitN*; ! Factor covariance (free)

MODEL CONSTRAINT:
NEW(SpearP); ! This is now equivalent to Spearman-Brown reliability for pos items
SpearP = (1*(PosLoad*3)**2) / ((1*(PosLoad*3)**2) + (PosError*3));

# R Model 7: Two Factors with Parallel Positive Items Only (loadings and error variances equal)
Syntax7 = " ! Note labels are in order of inclusion, not according to item name
# Define factor and request item factor Loadings all estimated
SitP =~ PosLoad*Sit2 + PosLoad*Sit4 + PosLoad*Sit6 # Pos items
SitN =~ L4*Sit1r + L5*Sit3r + L6*Sit5r # Neg items
# Item error variances all estimated
Sit2 ~~ PosError*Sit2; Sit4 ~~ PosError*Sit4; Sit6 ~~ PosError*Sit6 # Pos items
Sit1r ~~ E4*Sit1r; Sit3r ~~ E5*Sit3r; Sit5r ~~ E6*Sit5r # Neg items
# Calculate Spearman-Brown Reliability for Sum Score of Positive Factor:
SpearP := ((PosLoad*3)^2) / ( ((PosLoad*3)^2) + (PosError*3))
"
# Use MLR estimation like in Mplus, z-score latent factors (mean=0, SD=1)
Model17 = sem(model=Syntax7, data=Example4, estimator="MLR", mimic="mplus", std.lv=TRUE)
# Print solution: get fit, get effect size, STDYX solution
summary(object=Model17, fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)

print("LRT: Does the shared error variance for the positive items make fit worse?")
anova(Model17, Model16)

```

R lavaan output—notice that the positive standardized loadings are now equal, too:

Latent Variables:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP =~							
Sit2 (PsLd)		1.005	0.035	28.455	0.000	1.005	0.698
Sit4 (PsLd)		1.005	0.035	28.455	0.000	1.005	0.698
Sit6 (PsLd)		1.005	0.035	28.455	0.000	1.005	0.698
SitN =~							
Sit1r (L4)		1.325	0.048	27.817	0.000	1.325	0.759
Sit3r (L5)		1.347	0.044	30.622	0.000	1.347	0.845
Sit5r (L6)		1.011	0.055	18.407	0.000	1.011	0.589

Covariances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SitP ~~	SitN	0.581	0.040	14.582	0.000	0.581	0.581

Intercepts:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2		5.289	0.042	127.346	0.000	5.289	3.676
.Sit4		5.359	0.042	126.896	0.000	5.359	3.724
.Sit6		5.321	0.046	115.492	0.000	5.321	3.698
.Sit1r		4.547	0.053	86.474	0.000	4.547	2.604
.Sit3r		4.896	0.048	101.959	0.000	4.896	3.070
.Sit5r		4.860	0.052	94.060	0.000	4.860	2.832
SitP		0.000				0.000	0.000
SitN		0.000				0.000	0.000

Variances:

		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
.Sit2 (PsEr)		1.060	0.061	17.452	0.000	1.060	0.512
.Sit4 (PsEr)		1.060	0.061	17.452	0.000	1.060	0.512
.Sit6 (PsEr)		1.060	0.061	17.452	0.000	1.060	0.512

.Sit1r	(E4)	1.294	0.102	12.645	0.000	1.294	0.424
.Sit3r	(E5)	0.728	0.091	7.992	0.000	0.728	0.286
.Sit5r	(E6)	1.922	0.119	16.095	0.000	1.922	0.653
SitP		1.000				1.000	1.000
SitN		1.000				1.000	1.000

Defined Parameters:

	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
SpearP	0.741	0.020	36.909	0.000	0.741	0.741

[1] "LRT: Does the shared error variance for the positive items make fit worse?"

> anova(Model7, Model6)

Scaled Chi-Squared Difference Test (method = "satorra.bentler.2001")

lavaan NOTE:

The "Chisq" column contains standard test statistics, not the robust test that should be reported per model. A robust difference test is a function of two standard (not robust) statistics.

Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
Model6	10	22718	22803	38.7		
Model7	12	22754	22829	79.0	2	0.000037

Mplus model fit:

Number of Free Parameters	15
<hr/>	
Loglikelihood	
H0 Value	-11361.960
H0 Scaling Correction Factor	1.3443
for MLR	
H1 Value	-11322.435
H1 Scaling Correction Factor	1.4073
for MLR	

Does the assumption of parallel items hold for the positive items? How do we know?

Example results section describing these analyses...

(Note: You may borrow the phrasing contained in this example to describe various aspects of your analyses, but your own results sections will not mimic this example exactly—they should be customized to describe the what and the why of what you did, specifically, including theoretical motivation for any model modifications).

(Descriptive information for the sample and items would have already been given in the method section...)

The reliability and dimensionality of six items assessing forgiveness of situations was evaluated in a sample of 1,103 persons with confirmatory factor analyses using robust maximum likelihood estimation (MLR) in Mplus v. 8.8 (Muthén & Muthén, 1998–2017) and the R package lavaan (Rossell, 2012). All models were identified by setting any latent factor means to 0 and latent factor variances to 1, such that all item intercepts, item factor loadings, and item residual variances were then estimated. The six items utilized a seven-point response scale, and three items were reverse-coded prior to analysis such that higher values then indicated greater levels of situation forgiveness for all items. As reported in Table 1, model fit statistics include the obtained model χ^2 , its scaling factor (in which values different than 1.000 indicate deviations from multivariate normality), its degrees of freedom, and its p -value (in which non-significance is desirable for good fit), CFI (the Comparative Fit Index, in which values higher than .95 are desirable for good fit), and the RMSEA (the Root Mean Square Error of Approximation point estimate and 90% confidence interval, in which values lower than .06 are desirable for good fit). As reported in Table 2, nested model comparisons were conducted using the rescaled $-2\Delta LL$ with degrees of freedom equal to the rescaled difference in the number of parameters between models (i.e., a rescaled likelihood ratio test). The specific models examined are detailed below.

Although a one-factor model was initially posited to account for the pattern of covariance across these six items, it resulted in poor fit, as shown in Table 1. Although each item had a significant factor loading (with standardized loadings ranging from .509 to .778), a single latent factor did not adequately describe the pattern of relationship across these six items as initially hypothesized. Sources of local misfit were identified using the normalized residual covariance matrix, in which individual values were calculated as: (observed covariance – expected covariance) / SE of observed covariance. Relatively large positive residual covariances were observed among items 2, 4, and 6 (the positively-worded items), indicating that these items were more related than was predicted by the single-factor model.

Modification indices corroborated this pattern, further suggesting additional remaining relationships among the negatively-worded items as well.

The necessity of separate latent factors for the positively-worded and negatively-worded items was tested by specifying a two-factor model in which the positively-worded items 2, 4, and 6 indicated a *forgiveness* factor, and in which negatively-worded items 1, 3, and 5 indicated a *not unforgiveness* factor, and in which the two factors were allowed to covary. The two-factor model fit was acceptable by every criterion except the significant χ^2 , likely due to the large sample. In addition, the two-factor model fit significantly better than the one-factor model, as reported in Table 2, indicating that the estimated correlation between the two factors of .564 was significantly less than 1.000. Thus, the six items appeared to measure two separate but related constructs. Further examination of local fit via normalized residual covariances and modification indices yielded no interpretable remaining relationships, and thus this two-factor model was retained.

Table 3 provides the estimates and their standard errors for the item factor loadings, intercepts, and residual (error) variances from both the unstandardized and standardized solutions. All factor loadings and the factor covariance were statistically significant. As shown in Table 3, standardized loadings for the forgiveness factor items ranged from .625 to .759 (with R^2 values for the amount of item variance attributable to the factor ranging from .390 to .575), and standardized loadings for the not unforgiveness factor ranged from .588 to .846 (with R^2 values of .346 to .715), suggesting the factor loadings were practically significant as well. Omega model-based reliability was calculated for the sum scores of each factor as described in Brown (2015) as the squared sum of the factor loadings divided by the squared sum of the factor loadings plus the sum of the error variances plus twice the sum of the error covariances (although no error covariances were included here). Omega was .744 for the forgiveness factor and .775 for the not unforgiveness factor, suggesting marginal reliability for each of the three-item sum scores.

The resulting distribution of the factors was examined by requesting empirical Bayes estimates of the individual scores for each factor, as shown in Figure 1. Factor score standard errors for the forgiveness and not unforgiveness factors were .472 and .418, respectively. Factor score reliability was computed as the model-estimated factor variance (1.000) divided by that plus the squared factor score standard errors. Factor score reliability was .818 for the forgiveness factor and .851 for the not unforgiveness factor, suggesting acceptable reliability for each of the three-item factor scores.

In addition, Figure 2 shows the predicted response for each item as a linear function of the latent factor based on the estimated model parameters. As shown, the predicted item response goes above the highest response option just before a latent factor score of +2 (i.e., 2 SDs above the mean), resulting in a ceiling effect for both sets of factor scores, as also shown in Figure 1. In addition, for the not unforgiveness factor, the predicted item response goes below the lowest response option just before a latent factor score of -3 (i.e., 3 SDs below the mean), resulting in a floor effect for the not unforgiveness factor, as also shown in Figure 1.

The extent to which the items within each factor could be seen as exchangeable was then examined via an additional set of nested model comparisons, as reported in Table 1 (for fit) and Table 2 (for comparisons of fit). First, the assumption of tau-equivalence (i.e., true-score equivalence, equal discrimination across items) was examined by constraining the factor loadings to be equal within a factor. For the not unforgiveness factor, the tau-equivalent model fit was acceptable but was significantly worse than the original two-factor model fit (i.e., in which all loadings were estimated freely). For the forgiveness factor, however, the tau-equivalent model fit was acceptable and was not significantly worse than the original two-factor model fit. Thus, the assumption of tau-equivalence held for the forgiveness factor items only. Finally, the assumption of parallel items (i.e., equal factor loadings and equal residual variances, or equal reliability across items) was examined for the forgiveness factor items only, and the resulting model fit was acceptable but was significantly worse than the tau-equivalent forgiveness factor model fit. Thus, the assumption of parallel items did not hold for the forgiveness factor items. In summary, while the not unforgiveness factor items were not exchangeable, the forgiveness factor items were exchangeable with respect to their factor loadings only (i.e., equal discrimination, but not equal item residual variances or item reliability).

Tables would be built as seen in the excel workbook:

- Table 1 → “Model Fit Table 1” worksheet
- Table 2 → “MLR Comparisons Table 2” worksheet
- Table 3 → “Model Estimates Table 3” worksheet

Figures would be built as seen in this example:

- Figure 1 → Can be built in Mplus or ggplot2 in R
- Figure 2 → Can be built using “Factor Model Predictions” worksheet

References:

- Brown, T. A. (2015). *Confirmatory factor analysis for applied research* (2nd ed.). New York, NY: Guilford.
- Muthén, L. K., & Muthén, B.O. (1998–2017). *Mplus user's guide* (8th ed.). Los Angeles, CA: Muthén & Muthén.
- R Core Team (2017). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL <https://www.R-project.org/>.
- Rosseel, Y (2012). lavaan: An R Package for Structural Equation Modeling. *Journal of Statistical Software*, 48(2), 1-36. URL <http://www.jstatsoft.org/v48/i02/>.

Example 4 Continued: CFA of Forgiveness of Situations (N = 1103) using SAS MIXED

SAS Code to Read in Mplus Data:

```

* Import data from Mplus, becomes var1-var23 without names at top;
PROC IMPORT OUT=work.Situation DATAFILE= "&example.\Example4.csv" DBMS=CSV REPLACE;
    GETNAMES=NO; DATAROW=1; RUN;

* Rename variables, remove missing values;
DATA work.Situation; SET work.Situation;
    ARRAY old(23) var1-var23;
    ARRAY new(23) PersonID Self1 Self2r Self3 Self4r Self5 Self6r
        Other1r Other2 Other3r Other4 Other5r Other6
        Sit1r Sit2r Sit3r Sit4r Sit5r Sit6r
        Selfsub Othsub Sitsub HFSsum;
    DO i=1 TO 23; new(i)=old(i); IF new(i)=99999 THEN new(i)=.; END;
    DROP i var1-var23; RUN;

* Stack situation items into one column, as required by MIXED;
DATA work.SituationStacked; SET work.Situation;
    ARRAY aitem(6) Sit1r Sit2r Sit3r Sit4r Sit5r Sit6r;
    DO i=1 TO 6; itemnum=i; response=aitem(i); OUTPUT; END; DROP i; RUN;

```

Independence (Null) Baseline Model: Item means and variances, but NO covariances

```

TITLE "Independence (Null) CFA Model in MIXED";
PROC MIXED DATA=work.SituationStacked NOCLPRINT COVTEST IC NAMELEN=100 METHOD=ML;
    CLASS PersonID itemnum;
    MODEL response = itemnum / SOLUTION NOINT NOTEST;
    REPEATED itemnum / TYPE=TOEPPH(1) SUBJECT=PersonID R; RUN;

```

Estimated R Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	3.0493					
2		1.9028				
3			2.5431			
4				1.9672		
5					2.9451	
6						2.3412

TYPE=TOEPPH(1) predicts a diagonal matrix that would be the same as TYPE=UN(1).

Covariance Parameter Estimates					
Cov		Subject	Estimate	Standard	Z
Var(1)	PersonID	PersonID	3.0493	0.1298	23.48 <.0001
Var(2)	PersonID	PersonID	1.9028	0.08102	23.48 <.0001
Var(3)	PersonID	PersonID	2.5431	0.1083	23.48 <.0001
Var(4)	PersonID	PersonID	1.9672	0.08377	23.48 <.0001
Var(5)	PersonID	PersonID	2.9451	0.1254	23.48 <.0001
Var(6)	PersonID	PersonID	2.3412	0.09969	23.48 <.0001

The R matrix shows the unconditional variances per item—repeated in the next piece of output as Var(item). Note that this independence “null” model predicts NO covariances between items.

Information Criteria						
Neg2LogLike	Parms	AIC	AICC	HQIC	BIC	CAIC
24625.9	12	24649.9	24650.0	24672.6	24710.0	24722.0

Model fit is given as -2LL rather than LL (but otherwise is the same as given from Mplus).

Solution for Fixed Effects						
Effect	itemnum	Estimate	Error	DF	t Value	Pr > t
itemnum	1	4.5467	0.05258	5509	86.47 <.0001	
itemnum	2	5.2892	0.04153	5509	127.35 <.0001	
itemnum	3	4.8957	0.04802	5509	101.96 <.0001	
itemnum	4	5.3590	0.04223	5509	126.90 <.0001	
itemnum	5	4.8604	0.05167	5509	94.06 <.0001	
itemnum	6	5.3209	0.04607	5509	115.49 <.0001	

The fixed effects show the unconditional means per item.

Saturated (Unstructured) Baseline Model: Item means, variances, and covariances in original data

```
TITLE "Saturated (Unstructured) CFA Model in MIXED";
PROC MIXED DATA=work.SituationStacked NOCLPRINT COVTEST IC NAMELEN=100 METHOD=ML;
  CLASS PersonID itemnum;
  MODEL response = itemnum / SOLUTION NOINT NOTEST;
  REPEATED itemnum / TYPE=UN(6) SUBJECT=PersonID R RCORR; RUN;
```

TYPE=UN(6) predicts a fully-estimated R matrix with no constraints whatsoever.

Estimated R Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	3.0493	0.5772	1.8022	0.7339	1.3583	0.7946
2	0.5772	1.9028	0.6974	1.1029	0.6043	0.9652
3	1.8022	0.6974	2.5431	0.8244	1.3191	0.8676
4	0.7339	1.1029	0.8244	1.9672	0.6947	0.9618
5	1.3583	0.6043	1.3191	0.6947	2.9451	0.7982
6	0.7946	0.9652	0.8676	0.9618	0.7982	2.3412

The R matrix shows the unconditional variances and covariances for the items.

RCORR is the unconditional correlation matrix.

Note THIS IS THE DATA—the only discrepancies you'd see relative to descriptive statistics would be from missing data, as these are ML estimates (that assume MAR rather than MCAR as in listwise deletion).

Estimated R Correlation Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	1.0000	0.2396	0.6472	0.2997	0.4533	0.2974
2	0.2396	1.0000	0.3170	0.5700	0.2553	0.4573
3	0.6472	0.3170	1.0000	0.3686	0.4820	0.3555
4	0.2997	0.5700	0.3686	1.0000	0.2886	0.4482
5	0.4533	0.2553	0.4820	0.2886	1.0000	0.3040
6	0.2974	0.4573	0.3555	0.4482	0.3040	1.0000

Cov Parm	Subject	Covariance Parameter Estimates			
		Estimate	Standard	Z	Pr > Z
UN(1,1)	PersonID	3.0493	0.1298	23.48	<.0001
UN(2,1)	PersonID	0.5772	0.07458	7.74	<.0001
UN(2,2)	PersonID	1.9028	0.08102	23.48	<.0001
UN(3,1)	PersonID	1.8022	0.09988	18.04	<.0001
UN(3,2)	PersonID	0.6974	0.06948	10.04	<.0001
UN(3,3)	PersonID	2.5431	0.1083	23.48	<.0001
UN(4,1)	PersonID	0.7339	0.07699	9.53	<.0001
UN(4,2)	PersonID	1.1029	0.06705	16.45	<.0001
UN(4,3)	PersonID	0.8244	0.07178	11.49	<.0001
UN(4,4)	PersonID	1.9672	0.08377	23.48	<.0001
UN(5,1)	PersonID	1.3583	0.09907	13.71	<.0001
UN(5,2)	PersonID	0.6043	0.07356	8.21	<.0001
UN(5,3)	PersonID	1.3191	0.09148	14.42	<.0001
UN(5,4)	PersonID	0.6947	0.07543	9.21	<.0001
UN(5,5)	PersonID	2.9451	0.1254	23.48	<.0001
UN(6,1)	PersonID	0.7946	0.08393	9.47	<.0001
UN(6,2)	PersonID	0.9652	0.06988	13.81	<.0001
UN(6,3)	PersonID	0.8676	0.07798	11.13	<.0001
UN(6,4)	PersonID	0.9618	0.07081	13.58	<.0001
UN(6,5)	PersonID	0.7982	0.08264	9.66	<.0001
UN(6,6)	PersonID	2.3412	0.09969	23.48	<.0001

Information Criteria						
Neg2LogLike	Parms	AIC	AICC	HQIC	BIC	CAIC
22644.9	27	22698.9	22699.1	22750.0	22834.0	22861.0

Solution for Fixed Effects

Effect	itemnum	Standard				
		Estimate	Error	DF	t Value	Pr > t
itemnum	1	4.5467	0.05258	5509	86.47	<.0001
itemnum	2	5.2892	0.04153	5509	127.35	<.0001
itemnum	3	4.8957	0.04802	5509	101.96	<.0001
itemnum	4	5.3590	0.04223	5509	126.90	<.0001
itemnum	5	4.8604	0.05167	5509	94.06	<.0001
itemnum	6	5.3209	0.04607	5509	115.49	<.0001

The fixed effects again show the unconditional means per item.

Model 1. Single Factor with Fully Z-Scored Factor Model Identification (Factor Variance = 1, Factor Mean = 0, All Loadings and Intercepts Estimated)

```
TITLE "Single-Factor CFA Model (Factor Variance=1, Factor Mean=0) in MIXED";
PROC MIXED DATA=work.SituationStacked NOCLPRINT COVTEST IC NAMELEN=100 METHOD=ML;
  CLASS PersonID itemnum;
  MODEL response = itemnum / SOLUTION NOINT NOTEST;
  REPEATED itemnum / TYPE=FA(1) SUBJECT=PersonID R RCORR;
RUN;
```

TYPE=FA(1) creates the covariance matrix that would be predicted by a single-factor CFA model.

Estimated R Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	3.0493	0.8670	1.5313	0.9682	1.2626	1.0108
2	0.8670	1.9028	0.8716	0.5511	0.7187	0.5753
3	1.5313	0.8716	2.5431	0.9733	1.2692	1.0161
4	0.9682	0.5511	0.9733	1.9672	0.8025	0.6424
5	1.2626	0.7187	1.2692	0.8025	2.9451	0.8378
6	1.0108	0.5753	1.0161	0.6424	0.8378	2.3412

The R matrix shows the predicted variances and covariances for the items.

RCORR is the single-factor predicted correlation matrix.

THIS IS NO LONGER THE DATA. So the objective is to see how close this predicted covariance matrix is from the one given by the saturated model (which was the data).

Estimated R Correlation Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	1.0000	0.3600	0.5499	0.3953	0.4213	0.3783
2	0.3600	1.0000	0.3962	0.2848	0.3036	0.2726
3	0.5499	0.3962	1.0000	0.4351	0.4638	0.4164
4	0.3953	0.2848	0.4351	1.0000	0.3334	0.2994
5	0.4213	0.3036	0.4638	0.3334	1.0000	0.3191
6	0.3783	0.2726	0.4164	0.2994	0.3191	1.0000

THIS IS NO LONGER THE DATA. So the objective is to see how close this predicted covariance matrix is from the one given by the saturated model (which was the data).

Covariance Parameter Estimates					
Cov Parm	Subject	Estimate	Standard Error	Z Value	Pr > t
FA(1)	PersonID	1.5259	0.09440	16.16	<.0001
FA(2)	PersonID	1.4093	0.07096	19.86	<.0001
FA(3)	PersonID	1.0038	0.07755	12.94	<.0001
FA(4)	PersonID	1.3518	0.07071	19.12	<.0001
FA(5)	PersonID	1.8986	0.09312	20.39	<.0001
FA(6)	PersonID	1.6706	0.08330	20.05	<.0001
FA(1,1)	PersonID	1.2342	0.05332	23.15	<.0001
FA(2,1)	PersonID	0.7025	0.04720	14.88	<.0001
FA(3,1)	PersonID	1.2407	0.04783	25.94	<.0001
FA(4,1)	PersonID	0.7845	0.04679	16.76	<.0001
FA(5,1)	PersonID	1.0230	0.05202	19.67	<.0001
FA(6,1)	PersonID	0.8190	0.05019	16.32	<.0001

The FA(item) terms are the item residual variances. The FA(item, factor) terms are the item factor loadings.

So the total variance per item is given by: loading²(1) + error variance, as shown in the R matrix above.

$$\text{Item 1} = 1.2342^2(1) + 1.5259 = 3.0493$$

The covariance between items is given by their loadings multiplied together.

$$\text{Item 1 and 2 cov} = 1.2342 * 0.7025 = 0.8670$$

Information Criteria						
Neg2LogLike	Parms	AIC	AICC	HQIC	BIC	CAIC
23072.8	18	23108.8	23108.9	23142.9	23198.9	23216.9

Solution for Fixed Effects						
Effect	itemnum	Estimate	Standard Error	DF	t Value	Pr > t
itemnum	1	4.5467	0.05258	5509	86.47	<.0001
itemnum	2	5.2892	0.04153	5509	127.35	<.0001
itemnum	3	4.8957	0.04802	5509	101.96	<.0001
itemnum	4	5.3590	0.04223	5509	126.90	<.0001
itemnum	5	4.8604	0.05167	5509	94.06	<.0001
itemnum	6	5.3209	0.04607	5509	115.49	<.0001

The fixed effects now show the intercepts per item conditional on factor = 0 (which then are equal to the original item means).

Tau-Equivalent Items Single-Factor Model with Marker Item Factor Model Identification (Factor Variance = ?, Factor Mean = 0, All Loadings Equal at 1)

TITLE "Tau-Equivalent Items Single-Factor CFA Model (Factor Variance=1, Factor Mean=0) in MIXED";
PROC MIXED DATA=work.SituationStacked NOCLPRINT COVTEST IC NAMELEN=100 METHOD=ML;

```
CLASS PersonID itemnum;
MODEL response = itemnum / SOLUTION NOINT NOTEST;
RANDOM INTERCEPT / TYPE=UN SUBJECT=PersonID G V VCORR;
REPEATED itemnum / TYPE=TOEPH(1) SUBJECT=PersonID R; RUN;
```

A random intercept creates a constant source of covariance across all items.

Estimated R Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	2.0017					
2		1.1357				
3			1.4550			
4				1.0866		
5					2.0552	
6						1.4565

The **R** matrix shows the item residual variances.

The **G** matrix shows the variance due to the factor for all items.

V is the predicted covariance matrix from putting **G** and **R** back together, and **VCORR** is the predicted correlation matrix.

Aligning notation
In MLM = In CFA:

V = Sigma Σ
Z = Lambda Λ
G = Phi Φ (factor var-cov)
R = Psi Ψ (error var-cov)

Estimated G Matrix						
Row	Effect	ID	Col1	Col2	Col3	Col4
1	Intercept	1	0.9127			
Estimated V Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	2.9143	0.9127	0.9127	0.9127	0.9127	0.9127
2	0.9127	2.0483	0.9127	0.9127	0.9127	0.9127
3	0.9127	0.9127	2.3677	0.9127	0.9127	0.9127
4	0.9127	0.9127	0.9127	1.9993	0.9127	0.9127
5	0.9127	0.9127	0.9127	0.9127	2.9679	0.9127
6	0.9127	0.9127	0.9127	0.9127	0.9127	2.3691
Estimated V Correlation Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	1.0000	0.3735	0.3474	0.3781	0.3103	0.3473
2	0.3735	1.0000	0.4144	0.4510	0.3702	0.4143
3	0.3474	0.4144	1.0000	0.4195	0.3443	0.3853
4	0.3781	0.4510	0.4195	1.0000	0.3747	0.4194
5	0.3103	0.3702	0.3443	0.3747	1.0000	0.3442
6	0.3473	0.4143	0.3853	0.4194	0.3442	1.0000

Covariance Parameter Estimates

Cov Parm	Subject	Estimate	Error	Standard	Z
				Value	Pr > Z
UN(1,1)	PersonID	0.9127	0.04938	18.48	<.0001
Var(1)	PersonID	2.0017	0.09613	20.82	<.0001
Var(2)	PersonID	1.1357	0.05929	19.15	<.0001
Var(3)	PersonID	1.4550	0.07304	19.92	<.0001
Var(4)	PersonID	1.0866	0.05703	19.05	<.0001
Var(5)	PersonID	2.0552	0.09729	21.13	<.0001
Var(6)	PersonID	1.4565	0.07161	20.34	<.0001

Information Criteria

Neg2LogLike	Parms	AIC	AICC	HQIC	BIC	CAIC
23131.1	13	23157.1	23157.1	23181.7	23222.2	23235.2

Solution for Fixed Effects

Effect	itemnum	Estimate	Error	Standard	t Value	Pr > t
				DF		
itemnum	1	4.5467	0.05140	5510	88.45	<.0001
itemnum	2	5.2892	0.04309	5510	122.74	<.0001
itemnum	3	4.8957	0.04633	5510	105.67	<.0001
itemnum	4	5.3590	0.04257	5510	125.87	<.0001
itemnum	5	4.8604	0.05187	5510	93.70	<.0001
itemnum	6	5.3209	0.04635	5510	114.81	<.0001

The fixed effects still show the intercepts per item conditional on factor = 0 (which then are equal to the original item means).

**Parallel Items Single-Factor Model with Marker Item Factor Model Identification
(Factor Variance = ?, Factor Mean = 0, All Loadings = 1 and All Error Variances Equal)**

```
TITLE "Parallel Items Single-Factor CFA Model (Factor Variance=1, Factor Mean=0) in MIXED";
PROC MIXED DATA=work.SituationStacked NOCLPRINT COVTEST IC NAMELEN=100 METHOD=ML;
```

```
CLASS PersonID itemnum;
MODEL response = itemnum / SOLUTION NOINT NOTESE;
RANDOM INTERCEPT / TYPE=UN SUBJECT=PersonID G V VCORR;
REPEATED itemnum / TYPE=VC SUBJECT=PersonID R; RUN;
```

A random intercept creates a constant source of covariance across all items. A Type=VC R matrix means equal residual variance across items.

Estimated R Matrix for PersonID 1						
Row	Col1	Col2	Col3	Col4	Col5	Col6
1	1.5180					
2		1.5180				
3			1.5180			
4				1.5180		
5					1.5180	
6						1.5180

Estimated G Matrix

Person

Row	Effect	ID	Col1
1	Intercept	1	0.9401

The R matrix shows the item residual variances.

Row	Col1	Col2	Col3	Col4	Col5	Col6
1	2.4581	0.9401	0.9401	0.9401	0.9401	0.9401
2	0.9401	2.4581	0.9401	0.9401	0.9401	0.9401
3	0.9401	0.9401	2.4581	0.9401	0.9401	0.9401
4	0.9401	0.9401	0.9401	2.4581	0.9401	0.9401
5	0.9401	0.9401	0.9401	0.9401	2.4581	0.9401
6	0.9401	0.9401	0.9401	0.9401	0.9401	2.4581

The G matrix shows the variance due to the factor for all items.

V is the predicted covariance matrix from putting G and R back together, and VCORR is the predicted correlation matrix.

Row	Col1	Col2	Col3	Col4	Col5	Col6
1	1.0000	0.3825	0.3825	0.3825	0.3825	0.3825
2	0.3825	1.0000	0.3825	0.3825	0.3825	0.3825
3	0.3825	0.3825	1.0000	0.3825	0.3825	0.3825
4	0.3825	0.3825	0.3825	1.0000	0.3825	0.3825
5	0.3825	0.3825	0.3825	0.3825	1.0000	0.3825
6	0.3825	0.3825	0.3825	0.3825	0.3825	1.0000

This type of predicted marginal V covariance matrix has a special name in MLM: compound symmetry.

Covariance Parameter Estimates						
Cov Parm	Subject	Estimate	Standard		Z	
			Error	Value	Pr > Z	
UN(1,1)	PersonID	0.9401	0.05103	18.42	<.0001	
itemnum	PersonID	1.5180	0.02891	52.51	<.0001	

Information Criteria						
Neg2LogLike	Parms	AIC	AICC	HQIC	BIC	CAIC
23254.0	8	23270.0	23270.1	23285.2	23310.1	23318.1

Solution for Fixed Effects

Effect	itemnum	Standard					
		Estimate	Error	DF	t Value	Pr > t	
itemnum	1	4.5467	0.04721	5510	96.31	<.0001	
itemnum	2	5.2892	0.04721	5510	112.04	<.0001	
itemnum	3	4.8957	0.04721	5510	103.71	<.0001	
itemnum	4	5.3590	0.04721	5510	113.52	<.0001	
itemnum	5	4.8604	0.04721	5510	102.96	<.0001	
itemnum	6	5.3209	0.04721	5510	112.71	<.0001	

The fixed effects still show the intercepts per item conditional on factor = 0 (which then are equal to the original item means).

Unfortunately, multiple factor models in MIXED appear to be EFA models instead of CFA models, so no examples of two-factor models are given here. PROC CALIS can be used for CFA in SAS.