# **Confirmatory Factor Analysis**

Statsomat.com

23 December 2020

Warning: The automatic computation and interpretation delivered by the Statsomat should not completely replace the classical, made by humans graphical exploratory data analysis and statistical analysis. There may be data cases for which the Statsomat does not deliver the most optimal solution or output interpretation.

	-	•		. •	
Daci	~ 1:	240	rm.	<b>~+:</b>	a n
Basi		шо		au	OH

<b>Automati</b>	c static	tice fo	r tha	filo.
AIIIOMan	CSIAIIS	$110 \le 10$	rine	III 6.

File neuroticism.csv

Your selection for the encoding: Auto

Your selection for the decimal character: Auto

Observations (rows with at least one non-missing value): 250 Variables (columns with at least one non-missing value): 8

Variables considered continuous: 8

Variables considered continuous
N1
N2
N3
N4
E1
E2
E3
E4

Numerical variables considered binary or ordinal: 0

Character variables considered binary: 0

Character variables considered nominal and transformed to binary: 0

# **Model Syntax**

The following table describes the applied model equations in lavaan model syntax, either as entered by you in the text area (denoted by User=1) or established internally (User=0). The last column numbers the free parameters which are estimated.

Left hand side	Operator	Right hand side	User	Free parameter
NEUROT	=~	N1	1	0
NEUROT	=~	N2	1	1
NEUROT	=~	N3	1	2
NEUROT	=~	N4	1	3
EXTRAV	=~	E1	1	0
EXTRAV	=~	E2	1	4
EXTRAV	=~	E3	1	5
EXTRAV	=~	E4	1	6
N1	~~	N1	0	7
N2	~~	N2	0	8
N3	~~	N3	0	9
N4	~~	N4	0	10
E1	~~	E1	0	11
E2	~~	E2	0	12
E3	~~	E3	0	13
E4	~~	E4	0	14
NEUROT	~~	NEUROT	0	15
EXTRAV	~~	EXTRAV	0	16
NEUROT	~~	EXTRAV	0	17
N1	~1		0	18
N2	~1		0	19
N3	~1		0	20
N4	~1		0	21
E1	~1		0	22
E2	~1		0	23
E3	~1		0	24
E4	~1		0	25
NEUROT	~1		0	0
EXTRAV	~1		0	0

# **Assumptions**

Open issue

# **Model Settings**

# Outputs

# **Model Fit Summary**

lavaan 0.6--7 ended normally after 59 iterations

Estimator Optimization method Number of free parameters	ML NLMINB 25
Number of observations Number of missing patterns	250 1
Model Test User Model:	
Test statistic Degrees of freedom P-value (Chi-square)	13.285 19 0.824
Model Test Baseline Model:	
Test statistic Degrees of freedom P-value	1253.791 28 0.000
User Model versus Baseline Model:	
Comparative Fit Index (CFI) Tucker-Lewis Index (TLI)	1.000 1.007
Loglikelihood and Information Criteria:	
Loglikelihood user model (H0) Loglikelihood unrestricted model (H1)	-5748.501 -5741.858
Akaike (AIC) Bayesian (BIC) Sample-size adjusted Bayesian (BIC)	11547.002 11635.038 11555.786
Root Mean Square Error of Approximation:	
RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value RMSEA <= 0.05	0.000 0.000 0.034 0.990

# Standardized Root Mean Square Residual:

SRMR	0.018
------	-------

## Parameter Estimates:

Standard errors	Standard
Information	Observed
Observed information based on	Hessian

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

# Latent Variables:

NEUROT =~				
N1	1.000			
N2	0.942	0.052	18.146	0.000
N3	1.071	0.061	17.592	0.000
N4	0.997	0.052	19.217	0.000
EXTRAV =~				
E1	1.000			
E2	1.074	0.078	13.713	0.000
E2 E3	1.074 0.935	0.078 0.072	13.713 12.921	0.000
			201120	

## Covariances:

	Estimate	Std.Err	z-value	P(> z )
NEUROT ~~				
FYTRAV	-10 512	1 032	-5 442	0 000

# Intercepts:

rercepts.				
	Estimate	Std.Err	z-value	P(> z )
.N1	0.000	0.360	0.000	1.000
.N2	0.000	0.353	0.000	1.000
.N3	-0.000	0.404	-0.000	1.000
.N4	-0.000	0.360	-0.000	1.000
.E1	-0.000	0.379	-0.000	1.000
.E2	-0.000	0.391	-0.000	1.000
.E3	-0.000	0.360	-0.000	1.000
.E4	0.000	0.353	0.000	1.000
NEUROT	0.000			
EXTRAV	0.000			

## Variances:

	Estimate	Std.Err	z-value	P(> z )
.N1	7.025	0.914	7.688	0.000
.N2	8.746	1.002	8.731	0.000
.N3	11.760	1.336	8.799	0.000
.N4	7.188	0.921	7.802	0.000

.E1	12.802	1.580	8.104	0.000
.E2	11.671	1.603	7.283	0.000
.E3	12.192	1.464	8.327	0.000
.E4	15.972	1.668	9.574	0.000
NEUROT	25.335	2.902	8.730	0.000
EXTRAV	23.054	3.187	7.233	0.000

# **Completely Standardized Parameter Estimates**

.E1

Latent Variables:						
	est.std	Std.Err	z-value	P(> z )	ci.lower	ci.upper
NEUROT =~						
N1	0.885	0.018	49.253	0.000	0.850	0.920
N2	0.849	0.021	39.868	0.000	0.807	0.890
N3	0.844	0.022	38.674	0.000	0.801	0.886
N4	0.882	0.018	48.478	0.000	0.846	0.918
EXTRAV =~						
E1	0.802	0.029	27.214	0.000	0.744	0.860
E2	0.834	0.027	30.766	0.000	0.781	0.887
E3	0.789	0.031	25.839	0.000	0.730	0.849
E4	0.699	0.038	18.451	0.000	0.625	0.773
Covariances:						
	est.std	Std.Err	z-value	P(> z )	ci.lower	ci.upper
NEUROT ~~						
EXTRAV	-0.435	0.059	-7.397	0.000	-0.550	-0.320
Intercepts:						
	est.std	Std.Err	z-value	P(> z )	ci.lower	<pre>ci.upper</pre>
.N1	0.000	0.063	0.000	1.000	-0.124	0.124
.N2	0.000	0.063	0.000	1.000	-0.124	0.124
.N3	-0.000	0.063	-0.000	1.000	-0.124	0.124
.N4	-0.000	0.063	-0.000	1.000	-0.124	0.124
.E1	-0.000	0.063	-0.000	1.000	-0.124	0.124
.E2	0.000	0.063	0.000	1.000	-0.124	0.124
.E3	-0.000	0.063	-0.000	1.000	-0.124	0.124
.E4	0.000	0.063	0.000	1.000	-0.124	0.124
NEUROT	0.000				0.000	0.000
EXTRAV	0.000				0.000	0.000
Variances:						
	est.std	Std.Err	z-value		ci.lower	
.N1	0.217	0.032	6.829	0.000	0.155	0.279
.N2	0.280	0.036	7.753	0.000	0.209	0.351
.N3	0.288	0.037	7.832	0.000	0.216	0.360
.N4	0.222	0.032	6.921	0.000	0.159	0.285

7.556

0.000

0.264

0.450

0.047

0.357

.E2	0.305	0.045	6.746	0.000	0.216	0.393
.E3	0.377	0.048	7.810	0.000	0.282	0.471
.E4	0.511	0.053	9.654	0.000	0.408	0.615
NEUROT	1.000				1.000	1.000
EXTRAV	1.000				1.000	1.000

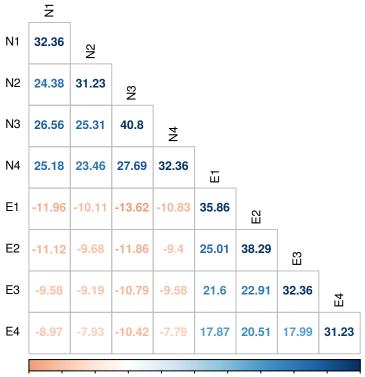
# **Communality**

Table 1: Communality

in which order is it?

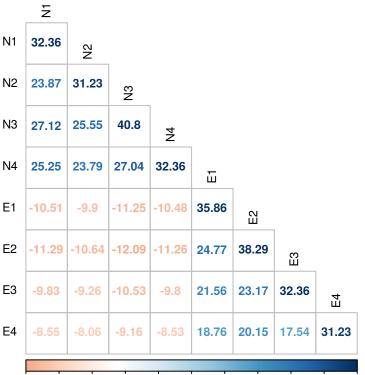
Variable	Communality
N1	0.78
N2	0.72
N3	0.71
N4	0.78
E1	0.64
E2	0.70
E3	0.62
E4	0.49

## **Observed Covariance Matrix**



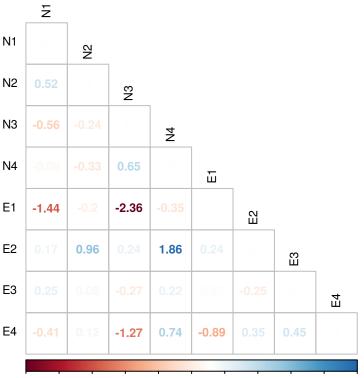
 $\textbf{-13.62-9.02-4.42\ 0.18\ 4.77\ 9.37\ 13.9718.5723.1627.7632.36}$ 

# Fitted Covariance Matrix



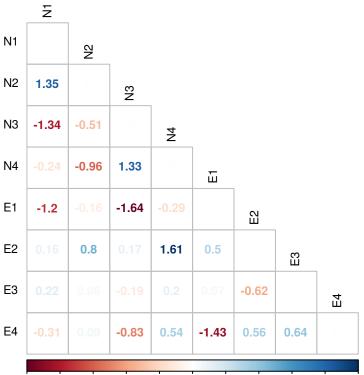
 $\hbox{-}12.09\hbox{-}7.65 \hskip1em \hbox{-}3.2 \hskip1em \hbox{1.24} \hskip1em \hbox{5.69} \hskip1em \hbox{10.13} \hskip1em \hbox{14.58} \hskip1em \hbox{19.022} \hskip1em \hbox{3.47} \hskip1em \hbox{27.91} \hskip1em \hbox{32.36}$ 

# **Residual Covariance Matrix**



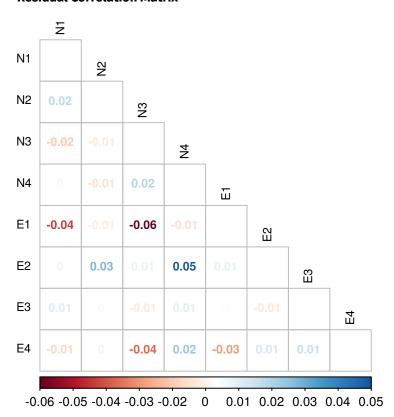
-2.36 -1.94 -1.52 -1.1 -0.67 -0.25 0.17 0.59 1.02 1.44 1.86

# **Standardized Residual Matrix**



-1.64 -1.31 -0.99 -0.67 -0.34 -0.02 0.31 0.63 0.96 1.28 1.61

# **Residual Correlation Matrix**



# **Modification Indices**

# define possible operators in the statsomet interface

Table 2: Modification Indices With Respect To (Residual) Correlation

Left	Operator	Right	Modification Index	Expected Parameter Change	Delta	Power	Decision
N3	~~	N4	3.507	1.820	0.1	0.051	(i)
N1	~~	N2	3.227	1.538	0.1	0.052	(i)
N1	~~	N3	2.747	-1.615	0.1	0.051	(i)
E1	~~	E4	1.983	-1.827	0.1	0.051	(i)
N4	~~	E2	1.276	0.906	0.1	0.052	(i)
N2	~~	N4	1.253	-0.956	0.1	0.052	(i)
N3	~~	E1	1.101	-1.028	0.1	0.051	(i)
N1	~~	E3	0.919	0.740	0.1	0.052	(i)
N4	~~	E3	0.838	-0.711	0.1	0.052	(i)
N2	~~	E1	0.658	0.689	0.1	0.052	(i)
N3	~~	E4	0.587	-0.785	0.1	0.051	(i)
E3	~~	E4	0.503	0.872	0.1	0.051	(i)
N1	~~	E2	0.500	-0.563	0.1	0.052	(i)
N4	~~	E4	0.480	0.584	0.1	0.052	(i)
E2	~~	E3	0.463	-1.037	0.1	0.050	(i)
E1	~~	E2	0.428	1.071	0.1	0.050	(i)
E2	~~	E4	0.412	0.873	0.1	0.051	(i)
N2	~~	N3	0.311	-0.525	0.1	0.051	(i)
N1	~~	E1	0.255	-0.405	0.1	0.052	(i)
N3	~~	E3	0.254	0.476	0.1	0.051	(i)
N2	~~	E3	0.222	-0.385	0.1	0.052	(i)
N1	~~	N4	0.121	-0.317	0.1	0.051	(i)
N3	~~	E2	0.107	0.319	0.1	0.051	(i)
N1	~~	E4	0.016	-0.107	0.1	0.052	(i)
E1	~~	E3	0.006	0.107	0.1	0.051	(i)
N2	~~	E2	0.004	-0.056	0.1	0.052	(i)
N2	~~	E4	0.002	0.041	0.1	0.051	(i)
N4	~~	E1	0.000	-0.005	0.1	0.052	(i)

Table 3: Modification Indices With Respect To Factor Loadings

Left	Operator	Right	Modification Index	Expected Parameter Change	Delta	Power	Decision
NEUROT	=~	E1	1.328	-0.073	0.4	1	(nm)
NEUROT	=~	E2	1.121	0.068	0.4	1	(nm)
EXTRAV	=~	N4	1.067	0.053	0.4	1	(nm)
EXTRAV	=~	N3	0.904	-0.059	0.4	1	(nm)
EXTRAV	=~	N1	0.273	-0.027	0.4	1	(nm)
EXTRAV	=~	N2	0.143	0.020	0.4	1	(nm)
NEUROT	=~	E3	0.014	0.007	0.4	1	(nm)
NEUROT	=~	E4	0.013	-0.007	0.4	1	(nm)

## Interpretation

## **Goodness of Fit Indices**

We consider some of the model fit indices from the Model Fit Summary section to check the goodness-of-fit of the model. To decide for an acceptable or non-acceptable model, we apply thresholds considered in the References: [@brown], [@kline].

#### **Model Test User Model**

The degrees of freedom are calculated as the number of known parameters minus the number of free parameters: 44 - 25 = 19. The 19 degrees of freedom indicate an over-identified model, fact which basically enables further analysis and interpretation.

The test statistic with the value 13.285 is called the Chi-square model fit index and represents the difference between summaries of the model-implied covariance matrix and the observed covariance matrix which is hypothesized and desirable to be zero. In general, if the p-value is larger than 0.05 then the test is not statistically significant at 5 % error, the hypothesis cannot be rejected, which would be in favour of the model.

In our case, the p-value is 0.824 suggesting an acceptable model fit.

#### **Model Test Baseline Model**

The test statistic with the value 1253.791 represents the difference between summaries of the baseline model (an alternative model-implied covariance matrix having zero covariances, i.e. a worst fitting model assuming independent variables) and the observed covariance matrix. The p-value of the test of a zero difference is <0.001 suggesting that the baseline model does not fit good to the data. This result is used indirectly in the construction of other model fit indices.

#### **Root Mean Square Error of Approximation:**

The Root Mean Square Error of Approximation (RMSEA) is a fit index based on the chi-square test statistic, which corrects for parsimony, i.e. overly complex models are penalized. RMSEA can be greater or equal than zero, with values close to zero suggesting an acceptable model fit.

In our case, the RMSEA is 0. The upper bound of the 90% confidence interval of the RMSEA is 0.034 and smaller than the threshold value 0.05, suggesting an excellent model fit.

#### **Standardized Root Mean Square Residual:**

The Standardized Root Mean Square Residual (SRMR) is a fit index derived from the residual correlation matrix with a range between zero and one with values close to zero suggesting an acceptable model fit.

In our case, the SRMR is 0.02 which is smaller than the threshold value 0.1 suggesting an acceptable model fit.

#### User Model versus Baseline Model

The Comparative Fit Index (CFI), evaluates the fit of the the model in relation to the worst-fitting baseline model described above. It ranges between zero and one, with values close to one suggesting good models (in the sense of departure from the baseline model).

In our case, the CFI is 1 which is greater or equal than the threshold value 0.95, suggesting a good model fit.

Similarly to the CFI, the Tucker-Lewis Index (TLI) evaluates the fit of the model in relation to the worst-fitting baseline model described above. Moreover, overly complex models are penalized. Values can range outside zero and one but the index is interpreted similarly to the CFI.

In our case, the TLI is 1.007 which is greater or equal than the threshold value 0.95, suggesting a good model fit.

#### **Summary of the Goodness of Fit Indices**

The Chi-square model fit index, the RMSEA and the TLI suggest an acceptable model fit. We tentatively assume an acceptable model fit and verify this assertion by considering further metrics.

#### Residuals

We analyze the residual matrices from the Outputs chapter. The residual covariance matrix represents the difference between the observed covariance matrix and the fitted model-implied covariance matrix. Large absolute values indicate local areas of misfit. However, the residuals are affected by the raw metric and are difficult to interpret more precisely.

A better interpretation allows the standardized residual matrix (residuals divided by their estimated asymptotic standard error) and the residual correlation matrix.

There are no variable pairs with standardized residuals which are larger or equal than the considered threshold 2.58 [@brown] or correlation residuals which are larger or equal than the considered threshold 0.1 [@kline]. Therefore, no relationships among the variables are substantially underestimated by the model. There are no variable pairs with standardized residuals which are smaller or equal than the considered threshold -2.58 [@brown] or correlation residuals which are smaller or equal than the considered threshold -0.1 [@kline]. Therefore, no relationships among the variables are substantially overestimated by the model.

#### **Modification Indices**

In the interpretation of the modification indices we rely mostly on [@brown] and [@mi]. We cite from [@brown]: "The modification index reflects an approximation of how much the overall model Chi² will decrease if the fixed or constrained parameter is freely estimated." In other words, if adding a line with a high modification index to the model, i.e. if adding a parameter, the overall goodness-of-fit may be improved. Nevertheless, this should be done only under certain conditions, described in the sequel.

We consider only modification indices greater or equal than 3.84 (which are statistically significant at 5% type I error). Next, we search only for modification indices which achieve a power of minimum 75% in detecting a (relevant) misspecification of at least 0.1 for (residual) correlations, respectively 0.4 for factor loadings. These are characterized in the decision column by the label "epc:m".

We remark that these conditions are not fulfilled for modification indices with respect to (residual) correlations. Therefore, there exist no significant and relevant modification indices with respect to (residual) correlations.

We remark that these conditions are not fulfilled for modification indices with respect to factor loadings. Therefore, there exist no significant and relevant modification indices with respect to factor loadings.

#### **Parameter Estimates**

#### **Factor Loadings**

We remark that the completely standardized factor loadings (section "Completely Standardized Parameter Estimates") are all statistically significant at 5% type I error. Moreover, in absolute value they are all greater than 0.4. This cutoff-value is considered in some CFA research areas a magnitude that is substantively meaningful [@brown]. Please consider also cutoff-values from your particular research area when interpreting the factor loadings. We summarize the interpretation of the completely standardized factor loadings in the next table:

Table 4: Check Completely Standardized Factor Loadings

Latent Variable	Observed Variable	Factor Loading <sup>1</sup>	P-Value	Significant? <sup>2</sup>	Relevance <sup>3</sup>	Direction	Check
NEUROT	N1	0.88	<0.001	Yes	***	Ok	Ok
NEUROT	N2	0.85	<0.001	Yes	***	Ok	Ok
NEUROT	N3	0.84	<0.001	Yes	***	Ok	Ok
NEUROT	N4	0.88	<0.001	Yes	***	Ok	Ok
EXTRAV	E1	0.80	<0.001	Yes	***	Ok	Ok
EXTRAV	E2	0.83	<0.001	Yes	***	Ok	Ok
EXTRAV	E3	0.79	<0.001	Yes	**	Ok	Ok
EXTRAV	E4	0.70	<0.001	Yes	**	Ok	Ok

<sup>&</sup>lt;sup>1</sup> The completely standardized factor loading can be interpreted as the correlation with the factor.

Next, we inspect the unstandardized factor loadings (section "Model Fit Summary"). We remark that the unstandardized factor loadings are all statistically significant at 5% type I error. Therefore, the significance test results for standardized and unstandardized factor loadings coincide (for non-marker variables). We summarize the interpretation of the unstandardized factor loadings in the next table(s):

Table 5: Check Unstandardized Factor Loadings

Latent Variable	Observed Variable	Factor Loading	P-Value	Significant? <sup>1</sup>	Direction
NEUROT	N1	1.00			Ok
NEUROT	N2	0.94	<0.001	Yes	Ok
NEUROT	N3	1.07	<0.001	Yes	Ok
NEUROT	N4	1.00	<0.001	Yes	Ok
EXTRAV	E1	1.00			Ok
EXTRAV	E2	1.07	<0.001	Yes	Ok
EXTRAV	E3	0.94	<0.001	Yes	Ok
EXTRAV	E4	0.81	<0.001	Yes	Ok

<sup>&</sup>lt;sup>1</sup> 5% type I error.

Table 6: Interpretation of Unstandardized Factor Loadings

Interpretation of Unstandardized Factor Loadings
N1 is marker variable for NEUROT
A 1-unit increase in NEUROT leads to a 0.94 -unit increase in the N2 $$
A 1-unit increase in NEUROT leads to a 1.07 -unit increase in the N3
A 1-unit increase in NEUROT leads to a 1.00 -unit increase in the N4 $$
E1 is marker variable for EXTRAV
A 1-unit increase in EXTRAV leads to a 1.07 -unit increase in the E2
A 1-unit increase in EXTRAV leads to a 0.94 -unit increase in the E3
A 1-unit increase in EXTRAV leads to a 0.81 -unit increase in the E4

<sup>&</sup>lt;sup>2</sup> 5% type I error.

<sup>&</sup>lt;sup>3</sup> Stars correspond to factor loadings cutoff-values: 0.4, 0.6, 0.8.

<sup>&</sup>lt;sup>4</sup> The observed variable is probably not related to latent factor.

<sup>&</sup>lt;sup>5</sup> Uncertain. The evidence is insufficient or the model is misspecified.

<sup>&</sup>lt;sup>6</sup> Uncertain. Significant but small(er) effect size.

#### **Factor Correlations**

As noted by [@brown], "the interpretability of the size and statistical significance of factor intercorrelations depends on the specific research context." Though, the largest estimated factor intercorrelation within the section "Completely Standardized Parameter Estimates" is -0.43 which we regard as a proof of a reasonable discriminant validity.

#### **Factor Reliability**

#### **Error Variances**

Table 7: Completely Standardized Error Variances and Communality

Observed Variable	Error Variance <sup>1</sup>	Communality <sup>23</sup>	P-Value	Significant Error Variance? <sup>4</sup>
E1	0.36	0.64	<0.001	Yes
E2	0.30	0.70	<0.001	Yes
E3	0.38	0.62	<0.001	Yes
E4	0.51	0.49	<0.001	Yes
N1	0.22	0.78	<0.001	Yes
N2	0.28	0.72	<0.001	Yes
N3	0.29	0.71	<0.001	Yes
N4	0.22	0.78	<0.001	Yes

<sup>&</sup>lt;sup>1</sup> Can be interpreted as proportion of unexplained variance by the latent factor(s) (%).

## Intercepts

In case of missing values and estimation via FIML, a meanstructure i.e. the intercepts of the observed variables are added to the model. The means of the latent factors are fixed to zero. Therefore, the estimated intercepts within the section "Model Fit Summary" are just the means of the observed variables.

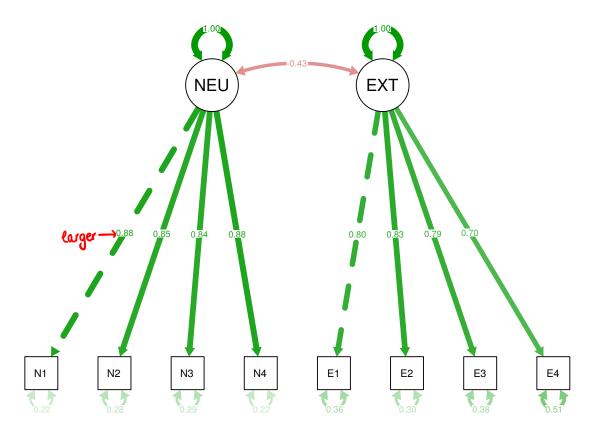
+ add: Is there a better model? - which?

<sup>&</sup>lt;sup>2</sup> Corresponds to the squared factor loading.

<sup>&</sup>lt;sup>3</sup> Can be interpreted as proportion of explained variance by the latent factor(s) (%).

<sup>&</sup>lt;sup>4</sup> 5% type I error. Typically significant since a large portion of variance is not explained by the latent variable.

# Path Diagram



# References