ESEM in R.

ESEM with lavaan is very simple. We first specify a model that looks just like the one for EFA in lavaan.

ESEM Model

The difference in the code is here. When fitting the model, we need to use rotation = geomin because geomin does not assume uncorrelated residuals. Like other oblique estimators, geomin allows latent factors to covary.

Note: In addition to changing the estimator, we should tell lavaan to change the epsilon value so that it overrides its default of 0.001. This is important: Changing epsilon allows to change the size of the cross-loadings or of the factor correlations. The lower the epsilon, the greater the inter-factor correlations, the lower the cross-loadings, and the less accurate the associations between the constructs (Morin et al., in press). To optimize the estimate of the inter-factor correlations, Morin et al. recommend epsilon = .5.

Important: By default, lavaan no longer allows changing the epsilon to values above .01, so the results will differ from those in Mplus and recommended values when using geomin. lavaan directly from GitHub may solve this, as their defaults are up to 1. To find lavaan defaults and how to change values: https://github.com/yrosseel/lavaan/blob/master/R/lav_options.R

We have missing data so we're telling lavaan to use FIML to estimate missing values and we're using MLR as our estimator.

```
## lavaan 0.6\text{--}20.2265 ended normally after 35 iterations ##
```

## ## ##	Estimator Optimization method Number of model parameters	ML NLMINB 37	
##	Row rank of the constraints matrix	2	
## ##	Potation mothod	CEOMIN ODI TOILE	
##	Rotation method Geomin epsilon	GEOMIN OBLIQUE 0.5	
##	Rotation algorithm (rstarts)	GPA (30)	
##	Standardized metric	TRUE	
##	Row weights	None	
##	0		
##		Used	Total
##	Number of observations	430	432
##	Number of missing patterns	40	
##			
##	Model Test User Model:		
##		Standard	Scaled
##	Test Statistic	79.099	60.953
##	Degrees of freedom	19	19
##	P-value (Chi-square)	0.000	0.000
##	8		1.298
##	Yuan-Bentler correction (Mplus vari	lant)	
	Model Test Baseline Model:		
##	model lest baseline model.		
##	Test statistic	1554.087	1043.454
##	Degrees of freedom	36	36
##	P-value	0.000	0.000
##	Scaling correction factor		1.489
##	<u> </u>		
##	User Model versus Baseline Model:		
##			
##	Comparative Fit Index (CFI)	0.960	0.958
##	Tucker-Lewis Index (TLI)	0.925	0.921
##			
##	Robust Comparative Fit Index (CFI)		0.965
##	Robust Tucker-Lewis Index (TLI)		0.933
##	Indibalihard and Information Chitamia		
##	Loglikelihood and Information Criteria:		
##	Loglikelihood user model (HO)	-5225.008	-5225.008
##	Scaling correction factor	0220.000	1.231
##	for the MLR correction		1.201
##	Loglikelihood unrestricted model (H1)	-5185.458	-5185.458
##	Scaling correction factor		1.254
##	for the MLR correction		
##			
##	Akaike (AIC)	10520.015	10520.015
##	Bayesian (BIC)	10662.248	10662.248
##	Sample-size adjusted Bayesian (SABIC)	10551.178	10551.178
##			
	Root Mean Square Error of Approximation	1:	
##	DWCHA	2 222	0 000
##	RMSEA	0.086	0.072

##	90 Percent conf	idence inte	rval - lo	wer	0.067	0.0	54
##	90 Percent conf	idence inte	rval - up	per	0.106	0.0	90
##	P-value H_0: RM	ISEA <= 0.05	0	_	0.001	0.0	21
##	P-value H_0: RM	ISEA >= 0.08	0		0.706	0.2	:34
##	_						
##							92
##	90 Percent conf	idence inte	rval - lo	wer		0.0	
##	90 Percent conf					0.1	
##	P-value H_0: Ro			r		0.0	
##	P-value H_0: Ro					0.7	
##	- · · · · · · · · · · · · · · · · · · ·						
	Standardized Root	: Mean Squar	e Residua	1:			
##		moun oquun	0 10001444				
##	SRMR				0.031	0.0	31
##	Diant				0.001	0.0	.01
	Parameter Estimat	05.					
##	Tarameter Ebtimat						
##	Standard errors	•			Sandwich		
##	Information bre				Observed		
##	Observed inform		on		Hessian		
##	upserved inform	lation based	. 011		пезатап		
	Latent Variables:						
	Latent variables:		C+ 3 E		P(> z)	C+1 1	רוי ויי
##	E1 - blask1	Estimate	Sta.EII	z-varue	P(> 2)	Std.lv	Std.all
##	F1 =~ block1	0.000	0 000	0.000	0 000	0 000	0 001
##	newhabit01	0.838	0.092		0.000	0.838	0.681
##	newhabit02	0.989	0.088	11.272	0.000	0.989	0.769
##	newhabit03	0.656	0.087		0.000	0.656	0.540
##	newhabit04	0.709	0.116	6.097		0.709	0.549
##	newhabit05	0.577	0.116	4.957		0.577	0.489
##	newhabit06	0.419	0.122	3.446	0.001	0.419	0.376
##	newhabit07	0.173	0.095	1.808		0.173	0.135
##	newhabit08	0.016	0.087			0.016	0.011
##	newhabit09	0.131	0.093	1.403	0.160	0.131	0.102
##	F2 =~ block1						
##	newhabit01	0.138	0.091	1.520	0.128		0.112
##	newhabit02	0.167	0.080	2.096	0.036	0.167	0.130
##	newhabit03	0.173	0.086	2.017	0.044	0.173	0.142
##	newhabit04	0.267	0.111	2.396	0.017	0.267	0.207
##	newhabit05	0.331	0.115	2.870	0.004	0.331	0.280
##	newhabit06	0.397	0.127	3.122	0.002	0.397	0.357
##	newhabit07	0.807	0.109	7.436	0.000	0.807	0.634
##	newhabit08	1.021	0.108	9.463	0.000	1.021	0.732
##	newhabit09	0.925	0.115	8.013	0.000	0.925	0.717
##							
##	Covariances:						
##		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	F1 ~~						
##	F2	0.553	0.038	14.510	0.000	0.553	0.553
##							
##	Intercepts:						
##		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	.newhabit01	3.405	0.062	55.325	0.000	3.405	2.768
##	.newhabit02	3.230	0.063	51.352	0.000	3.230	2.509
##	.newhabit03	2.907	0.059	49.072	0.000	2.907	2.393

##	.newhabit04	3.083	0.063	49.088	0.000	3.083	2.388
##	.newhabit05	3.263	0.057	56.876	0.000	3.263	2.764
##	.newhabit06	3.533	0.055	64.663	0.000	3.533	3.175
##	.newhabit07	3.515	0.064	54.618	0.000	3.515	2.759
##	.newhabit08	3.275	0.070	46.550	0.000	3.275	2.347
##	.newhabit09	3.213	0.064	50.490	0.000	3.213	2.492
##							
##	Variances:						
##		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	.newhabit01	0.664	0.094	7.066	0.000	0.664	0.439
##	.newhabit02	0.468	0.085	5.525	0.000	0.468	0.282
##	.newhabit03	0.891	0.092	9.685	0.000	0.891	0.603
##	.newhabit04	0.885	0.095	9.353	0.000	0.885	0.531
##	.newhabit05	0.740	0.068	10.945	0.000	0.740	0.531
##	.newhabit06	0.721	0.072	10.024	0.000	0.721	0.583
##	.newhabit07	0.788	0.100	7.857	0.000	0.788	0.485
##	.newhabit08	0.886	0.139	6.396	0.000	0.886	0.455
##	.newhabit09	0.655	0.130	5.041	0.000	0.655	0.394
##	F1	1.000				1.000	1.000
##	F2	1.000				1.000	1.000
##							
##	R-Square:						
##		Estimate					
##	newhabit01	0.561					
##	newhabit02	0.718					
##	newhabit03	0.397					
##	newhabit04	0.469					
##	newhabit05	0.469					
##	newhabit06	0.417					
##	newhabit07	0.515					
##	newhabit08	0.545					
##	newhabit09	0.606					

Notice that the correlation between the two latent variables is much lower in the ESEM model (r = .55) than in the EFA model (r = .71), and in the CFA model (r = .78) which makes sense given our use of geomin and specification of the value of epsilon. In the context of ESEM, it also makes sense that correlations would be lower in the ESEM model, as not allowing cross-loadings leads to inflation of the correlation between latent factors.

Cross-loadings are fairly small in the ESEM solution. They range from .01-.14 for the first factor (and are all non-significant), and .11-36 for the second factor (all but one are significant). The factors are fairly well-defined in the ESEM solution: the loadings on f1 range from .38-.77 (M = .57), and on f2 range from .63-.73 (M = .69). Model fit is good, but it is poorer than that of the Bifactor CFA model when we consider the TLI and RMSEA (which tax low parsimony).

Bifactor ESEM

To estimate a bifactor ESEM model, just add a general factor to the model with the same specifications as the two specific factors.

```
# these are exploratory blocks (you can give them any name), you'll need to specify them
biesem_model <- 'efa("block1")*F1 =~ newhabit01 + newhabit02 + newhabit03 + newhabit04 +</pre>
```

```
newhabit05 + newhabit06 + newhabit07 + newhabit08 +
newhabit09

efa("block1")*F2 =~ newhabit01 + newhabit02 + newhabit03 + newhabit04 +
newhabit05 + newhabit06 + newhabit07 + newhabit08 +
newhabit09

efa("block1")*g =~ newhabit01 + newhabit02 + newhabit03 + newhabit04 +
newhabit05 + newhabit06 + newhabit07 + newhabit08 +
newhabit09'
```

You'll need to change the rotation type to "bigeomin", which is one of the recommended rotations for bifactor ESEM (also what we're using in Mplus demo).

Compare the output with the one from Mplus. You'll notice that the standardized solution here is very similar to that in Mplus (under STDYX), but not exactly the same. This is likely because Mplus is using BI-GEOMIN orthogonal, in which the specific factors are uncorrelated with the general factor and with each other. Notice that lavaan is using "BIGEOMIN OBLIQUE" and that covariances/correlations are indeed non-zero (even if very small). Setting the factors to be orthogonal does not change lavaan's default to let the LV correlate.

```
lavaan::summary(biesemfit,
    fit.measures = TRUE,
    standardized = TRUE,
    rsquare = TRUE)
```

```
## lavaan 0.6-20.2265 ended normally after 49 iterations
##
##
     Estimator
                                                          ML
##
     Optimization method
                                                     NLMINB
     Number of model parameters
##
                                                          48
     Row rank of the constraints matrix
##
##
                                           BIGEOMIN OBLIQUE
##
     Rotation method
     Rotation algorithm (rstarts)
                                                   GPA (30)
##
     Standardized metric
                                                        TRUE
##
##
     Row weights
                                                       None
##
##
                                                        Used
                                                                   Total
##
     Number of observations
                                                         430
                                                                     432
##
     Number of missing patterns
                                                          40
##
## Model Test User Model:
                                                   Standard
                                                                  Scaled
##
##
     Test Statistic
                                                     34.932
                                                                  44.017
##
     Degrees of freedom
                                                                      12
                                                          12
     P-value (Chi-square)
                                                      0.000
                                                                   0.000
```

##	Scaling correction factor		0.794	
##	Scaling correction factor Yuan-Bentler correction (Mplus variant)		0.794	
##	ruan benefici coffeetion (nprus variant)			
	Model Test Baseline Model:			
##	Hodel Test Baseline Hodel.			
##	Test statistic	1554.087	1043.454	
##	Degrees of freedom	36	36	
##	P-value	0.000	0.000	
##	Scaling correction factor	0.000	1.489	
##	bearing correction factor		1.403	
	User Model versus Baseline Model:			
##	ober moder verbub baberine moder.			
##	Comparative Fit Index (CFI)	0.985	0.968	
##	Tucker-Lewis Index (TLI)	0.955	0.905	
##	rucker hewib mack (1817)	0.000	0.000	
##	Robust Comparative Fit Index (CFI)		0.997	
##	Robust Tucker-Lewis Index (TLI)		0.991	
##	Nobabi Tacker Bewis Index (IBI)		0.001	
	Loglikelihood and Information Criteria:			
##				
##	Loglikelihood user model (HO)	-5202.924	-5202.924	
##	Scaling correction factor	02021021	1.386	
##	for the MLR correction			
##	Loglikelihood unrestricted model (H1)	-5185.458	-5185.458	
##	Scaling correction factor		1.254	
##	for the MLR correction			
##				
##	Akaike (AIC)	10489.848	10489.848	
##	Bayesian (BIC)	10660.527	10660.527	
##	Sample-size adjusted Bayesian (SABIC)	10527.244	10527.244	
##				
##				
##	Root Mean Square Error of Approximation:			
##	Root Mean Square Error of Approximation:			
	Root Mean Square Error of Approximation:	0.067	0.079	
##	-	0.067 0.042	0.079 0.052	
## ##	RMSEA			
## ## ##	RMSEA 90 Percent confidence interval - lower	0.042	0.052	
## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper	0.042 0.093	0.052 0.108	
## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050	0.042 0.093 0.128	0.052 0.108 0.040	
## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA	0.042 0.093 0.128	0.052 0.108 0.040 0.506	
## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025	
## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025 0.165	
## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114	
## ## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025 0.165	
## ## ## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114	
## ## ## ## ## ## ## ## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114	
## ## ## ## ## ## ## ## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080 Standardized Root Mean Square Residual:	0.042 0.093 0.128 0.217	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114 0.703	
## ## ## ## ## ## ## ## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080	0.042 0.093 0.128	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114	
## ## ## ## ## ## ## ## ## ## ## ## ##	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080 Standardized Root Mean Square Residual: SRMR	0.042 0.093 0.128 0.217	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114 0.703	
## # # # # # # # # # # # # # # # # # #	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080 Standardized Root Mean Square Residual:	0.042 0.093 0.128 0.217	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114 0.703	
## # # # # # # # # # # # # # # # # # #	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080 Standardized Root Mean Square Residual: SRMR Parameter Estimates:	0.042 0.093 0.128 0.217	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114 0.703	
## # # # # # # # # # # # # # # # # # #	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080 Standardized Root Mean Square Residual: SRMR Parameter Estimates: Standard errors	0.042 0.093 0.128 0.217	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114 0.703	
## # # # # # # # # # # # # # # # # # #	RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: RMSEA <= 0.050 P-value H_0: RMSEA >= 0.080 Robust RMSEA 90 Percent confidence interval - lower 90 Percent confidence interval - upper P-value H_0: Robust RMSEA <= 0.050 P-value H_0: Robust RMSEA >= 0.080 Standardized Root Mean Square Residual: SRMR Parameter Estimates:	0.042 0.093 0.128 0.217	0.052 0.108 0.040 0.506 0.096 0.025 0.165 0.114 0.703	

##							
	Latent Variables:						
##		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	F1 =~ block1						
##	newhabit01	0.853	0.060	14.198	0.000	0.853	0.694
##	newhabit02	1.054	0.044	23.823	0.000	1.054	0.819
##	newhabit03	0.721	0.062	11.626	0.000	0.721	0.594
##	newhabit04	0.856	0.054	15.750	0.000	0.856	0.663
##	newhabit05	0.872	0.127	6.866	0.000	0.872	0.739
##	newhabit06	0.719	0.072	10.008	0.000	0.719	0.646
##	newhabit07	0.843	0.065	12.899	0.000	0.843	0.662
##	newhabit08	0.895	0.068	13.225	0.000	0.895	0.642
## ##	newhabit09 F2 =~ block1	0.921	0.060	15.260	0.000	0.921	0.714
##	newhabit01	-0.273	0.078	-3.498	0.000	-0.273	-0.222
##	newhabit01	-0.498	0.321	-1.550	0.121	-0.498	-0.387
##	newhabit03	-0.158	0.021	-1.651	0.099	-0.158	-0.130
##	newhabit04	-0.187	0.089	-2.098	0.036	-0.187	-0.145
##	newhabit05	0.012	0.233	0.052	0.958	0.012	0.010
##	newhabit06	0.075	0.131	0.571	0.568	0.075	0.067
##	newhabit07	0.318	0.131	2.427	0.015	0.318	0.250
##	newhabit08	0.463	0.213	2.177	0.029	0.463	0.332
##	newhabit09	0.372	0.177	2.099	0.036	0.372	0.289
##	g =~ block1						
##	newhabit01	0.038	0.276	0.137	0.891	0.038	0.031
##	newhabit02	-0.208	0.092	-2.251	0.024	-0.208	-0.161
##	newhabit03	0.015	0.254	0.059	0.953	0.015	0.012
##	newhabit04	-0.120	0.211	-0.568	0.570	-0.120	-0.093
##	newhabit05	0.671	0.766	0.876	0.381	0.671	0.568
##	newhabit06	0.205	0.274	0.749	0.454	0.205	0.184
##	newhabit07	-0.098	0.088	-1.120	0.263	-0.098	-0.077
##	newhabit08	-0.175	0.277	-0.631	0.528	-0.175	-0.125
## ##	newhabit09	-0.210	0.327	-0.643	0.521	-0.210	-0.163
##	Covariances:						
##	covariances.	Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	F1 ~~	Lboimacc	Dournin	Z varuo	1 (7 21)	Dou.iv	Dou.ull
##	F2	0.000	0.000	6.993	0.000	0.000	0.000
##	g	0.000	0.000	1.323	0.186	0.000	0.000
##	F2 ~~						
##	g	-0.052	0.087	-0.601	0.548	-0.052	-0.052
##							
##	Intercepts:						
##		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	.newhabit01	3.406	0.061	55.406	0.000	3.406	2.770
##	.newhabit02	3.229	0.063	51.245	0.000	3.229	2.510
##	.newhabit03	2.907	0.059	49.009	0.000	2.907	2.392
##	.newhabit04	3.082	0.063	48.941	0.000	3.082	2.386
##	.newhabit05	3.266	0.058	56.795	0.000	3.266	2.767
##	.newhabit06	3.537	0.055	64.780	0.000	3.537	3.176
## ##	.newhabit07 .newhabit08	3.515 3.273	0.064 0.070	54.571 46.511	0.000 0.000	3.515 3.273	2.759 2.346
##	.newhabit09	3.213	0.070	50.476	0.000	3.213	2.492
##	. IIOWIIGDI 609	0.210	0.004	00.710	3.000	0.210	2.432

##	Variances:						
##		Estimate	Std.Err	z-value	P(> z)	Std.lv	Std.all
##	.newhabit01	0.707	0.084	8.444	0.000	0.707	0.468
##	.newhabit02	0.265	0.306	0.866	0.387	0.265	0.160
##	.newhabit03	0.931	0.097	9.609	0.000	0.931	0.631
##	.newhabit04	0.889	0.108	8.225	0.000	0.889	0.533
##	.newhabit05	0.184	1.248	0.147	0.883	0.184	0.132
##	.newhabit06	0.676	0.193	3.503	0.000	0.676	0.546
##	.newhabit07	0.798	0.103	7.787	0.000	0.798	0.492
##	.newhabit08	0.892	0.134	6.680	0.000	0.892	0.458
##	.newhabit09	0.623	0.154	4.045	0.000	0.623	0.375
##	F1	1.000				1.000	1.000
##	F2	1.000				1.000	1.000
##	g	1.000				1.000	1.000
##							
##	R-Square:						
##		Estimate					
##	newhabit01	0.532					
##	newhabit02	0.840					
##	newhabit03	0.369					
##	newhabit04	0.467					
##	newhabit05	0.868					
##	newhabit06	0.454					
##	newhabit07	0.508					
##	newhabit08	0.542					
##	newhabit09	0.625					

Model	df	CFI	TLI	RMSEA	SRMR
CFA: One factor	36	0.90	0.87	0.09	0.05
CFA: Two correlated factors	26	0.96	0.94	0.08	0.04
CFA: One factor plus method factor	24	0.96	0.94	0.08	0.04
CFA: second order	27	0.95	0.93	0.09	0.13
Bifactor	18	0.98	0.97	0.05	0.02
ESEM-geomin	19	0.97	0.93	0.07	0.03

So, is the bifactor model the best fit? Does this mean we can treat this measure as a single score? Bifactor models can accommodate patterns of association in the data that are not accounted for when we use correlated-factor models. For example, in this measure, it would be unreasonable to assume that the last three items are completely unrelated to the first six - there's conceptual overlap between them and other items - but the correlated-factors model does not account for this, resulting in lower fit relative to the bifactor and ESEM models.

This was a simple example, as what we're talking about (wording effects) is likely construct-irrelevant variance, or a methodological artifact. The purpose of this exercise - including both scripts - was to demonstrate the use of these tools to examine measure dimensionality. In this case, to determine whether it is tenable to use this measure as a composite index in an analysis.

Importantly, how one treats the structure of a construct depends on more than fit. As you've read on Chapter 18 of the Handbook, better fit does **not** mean that the model is an accurate representation of the measurement structure for everyone (or even for most people in the sample) or an accurate representation of the "true" latent structure itself. This doesn't mean that these models aren't useful, but that they can help understand the sources of item variance and inform whether a measurement structure is tenable and interpretable/valid. These decisions must be made with justifications beyond statistical tests and fit coefficients.