# **Cross-lagged Panel Model of Positive and Negative Social Exchanges**

Note: I use listwise deletion and robust estimation here (Satorra-Bentler corrections) for convenience, specified as MLM in Mplus and lavaan. With attrition, the MAR assumption deserves additional consideration, but may be a reasonable or more reasonable approach with this example (see Newsom, 2015, for a more in-depth discussion of the topic of missing data with longitudinal models).

title: Cross-lag panel model of positive and negative exchanges;							
<pre>data: file=c:\jason\mplus\semclass\long1.dat; format=4f9.6;     listwise=on;</pre>							
<pre>variable: names = pos posf neg negf; missing=blank;</pre>							
	.s: type=o nator=mlm;	general;					
	negf on ne posf on ne neg with p negf with	eg pos; pos;					
output:	stdyx;						
(output e	excerpts)						
Number of	observati	lons				151	
TESTS OF	MODEL FIT						
Chi-Squar	e Test of	Model Fit					
	P-Value	of Freedom Correction Fa 1	actor	0.000* 0 0.0000 Undefined	-		
* The chi-square value for MLM, MLMV, MLR, ULSMV, WLSM and WLSMV cannot be used for chi-square difference tests. MLM, MLR and WLSM chi-square difference testing is described in the Mplus Technical Appendices at www.statmodel.com. See chi-square difference testing in the index of the Mplus User's Guide.							
CFI/TLI							
	CFI TLI			1.000 1.000			
RMSEA (Root Mean Square Error Of Approximation)							
	Estimate			0.000			
SRMR (Standardized Root Mean Square Residual)							
	Value			0.000			
MODEL RESULTS							
		Estimate	S.E.	Est./S.E.	Two-Tailed P-Value		
NEGF NEG POS	ON	0.630 -0.075	0.093 0.054	6.740 -1.395	0.000 0.163		
POSF NEG POS	ON	-0.139 0.532	0.103 0.076	-1.353 6.968	0.176		

NEG POS	WITH	-0.191	0.051	-3.734	0.000
NEGF POSF	WITH	-0.069	0.021	-3.233	0.001
Means POS NEG		2.778 0.629	0.062 0.044	44.634 14.301	0.000 0.000
Intercep POSF NEGF	ts	1.476 0.515	0.265 0.185	5.562 2.784	0.000 0.005
Variance POS NEG	S	0.585 0.292	0.079 0.053	7.410 5.517	0.000
Residual POSF NEGF	Variances	0.199 0.178	0.027 0.026	7.461 6.933	0.000

STANDARDIZED MODEL RESULTS

#### STDYX Standardization

		Estimate	S.E.	Est./S.E.	Two-Tailed P-Value
NEGF NEG POS	ON	0.607 -0.102	0.068 0.073	8.910 -1.410	0.000 0.159
POSF NEG POS	ON	-0.119 0.644	0.085 0.064	-1.398 10.119	0.162
NEG POS	WITH	-0.463	0.088	-5.256	0.000
NEGF POSF	WITH	-0.366	0.092	-3.976	0.000
Means POS NEG		3.632 1.164	0.295 0.087	12.311 13.426	0.000 0.000
Intercep POSF NEGF	ts	2.338 0.917	0.508 0.334	4.606 2.750	0.000 0.006
Variance POS NEG	S	1.000	0.000	999.000 999.000	999.000 999.000
Residual POSF NEGF	Variances	0.500 0.564	0.061 0.068	8.252 8.311	0.000 0.000

### R-SQUARE

Observed Variable	Estimate	S.E.	Est./S.E.	Two-Tailed P-Value
POSF	0.500	0.061	8.253	0.000
NEGF	0.436	0.068	6.425	

#### lavaan

```
mydata <- mydata[complete.cases(mydata), ]
library(psych)
</pre>
>
>
  #describe(mydata)
>
>
>
> library(lessR)
> model =
  posf ~ pos + neg
negf ~ pos + neg
+
/ #note: including the correlations between the exogenous variables
/ # and endogenous disturbances causes an error, so omit (still estimated by default)
/ fit = sem(model, data = mydata, mimic="Mplus", missing = 'listwise', estimator="MLM")
/ summary(fit,fit.measures=TRUE, rsquare=TRUE, standardized=TRUE)
lavaan 0.6.15 ended normally after 32 iterations
   Estimator
                                                                       ML
  Optimization method
                                                                  NLMINB
  Number of model parameters
                                                                         9
  Number of observations
                                                                     151
Model Test User Model:
                                                               Standard
                                                                                  Scaled
  Test Statistic
                                                                   0.000
                                                                                   0.000
  Degrees of freedom
                                                                        0
                                                                                         0
Model Test Baseline Model:
  Test statistic
                                                                212.861
                                                                                 139.021
  Degrees of freedom
P-value
                                                                   0.000
                                                                                   0.000
  Scaling correction factor
                                                                                   1.531
User Model versus Baseline Model:
                                                                                   1.000
   Comparative Fit Index (CFI)
                                                                   1.000
  Tucker-Lewis Index (TLI)
                                                                   1.000
                                                                                   1.000
   Robust Comparative Fit Index (CFI)
                                                                                        NA
  Robust Tucker-Lewis Index (TLI)
                                                                                        NA
Loglikelihood and Information Criteria:
   Loglikelihood user model (HO)
                                                               -165.435
                                                                               -165.435
  Loglikelihood unrestricted model (H1)
                                                               -165.435
                                                                               -165.435
                                                                348.870
376.026
   Akaike (AIC)
                                                                                 348.870
  Bayesian (BIC)
Sample-size adjusted Bayesian (SABIC)
                                                                                 376.026
                                                                347.542
                                                                                 347.542
Root Mean Square Error of Approximation:
   RMSEA
                                                                   0.000
                                                                                        NA
  90 Percent confidence interval - lower
90 Percent confidence interval - upper
                                                                   0.000
                                                                                        NA
                                                                   0.000
                                                                                        NA
  P-value H_0: RMSEA <= 0.050
P-value H_0: RMSEA >= 0.080
                                                                       NA
                                                                                        NA
                                                                       NA
                                                                                        NA
                                                                                   0.000
  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
                                                                                   0.000
                                                                                   0.000
                                                                                       NΔ
                                                                                       NA
Standardized Root Mean Square Residual:
                                                                   0.000
                                                                                   0.000
  SRMR
Parameter Estimates:
   Standard errors
                                                            Robust.sem
  Information
                                                               Expected
  Information saturated (h1) model
                                                            Structured
Regressions:
                          Estimate Std.Err z-value P(>|z|)
                                                                              Std.lv Std.all
   posf ~
                              0.532
                                          0.076
                                                       6.968
                                                                   0.000
                                                                               0.532
                                                                                            0.644
     pos
```

Newsom Psy 523/623 Structural Equation Modeling, Spring 2023

neg	-0.139	0.103	-1.353	0.176	-0.139	-0.119
negf~	-0.139	0.105	-1.333	0.170	-0.139	-0.119
pos	-0.075	0.054	-1.395	0.163	-0.075	-0.102
neg	0.630	0.093	6.740	0.000	0.630	0.607
Covariances:						
nacf	Estimate	Std.Err	z-value	P(> z )	Std.lv	Std.all
.posf ~~ .negf	-0.069	0.021	-3.233	0.001	-0.069	-0.366
Intercepts:						
	Estimate	Std.Err		P(> z )	Std.lv	Std.all
.posf .negf	1.476 0.515	0.265 0.185	5.561 2.785	0.000 0.005	$1.476 \\ 0.515$	2.338 0.917
5	01020	0.200		01000	01020	0101
Variances:	Estimate	Std.Err	z-value	P(> z )	Std.lv	Std.all
.posf	0.199	0.027	7.461	0.000	0.199	0.500
negf	0.178	0.026	6.933	0.000	0.178	0.564
R-Square:	<b>_</b> . • .					
posf	Estimate 0.500					
negf	0.436					

The output only shows the correlation between endogenous disturbances, but it is still estimating the correlation between exogenous variables. lavaan and Mplus are sometimes quirky about showing exogenous correlations with measured variables.<sup>1</sup> How do I know they are being estimated? Because the df is equal to 0, so I know that all possible relations are being estimated in this model.

## Write-up

A cross-lagged panel model was tested to investigate the longitudinal effects of positive and negative social exchanges over a thirteen-week interval. The model was just identified, so there was no information about fit. Both negative and positive social exchange measures were highly stable over the thirteen weeks,  $\beta = .630$ , SE = .093,  $\beta^* = .607$ , p < 001, and  $\beta = .532$ , SE = .076,  $\beta^* = .644$ , p < .001, respectively. Although the two measures were significantly negatively correlated at baseline,  $\psi^* = .463$ , p < .001, and the disturbances were significantly correlated at follow-up,  $\psi^* = .366$ , p < .001, neither cross-lagged effect was significant. Positive exchanges at baseline did not significantly predict negative exchanges at follow-up,  $\beta = -.075$ , SE = .054,  $\beta^* = .102$ , ns, and negative social exchanges at baseline did not significantly predictive positive social exchanges at follow-up,  $\beta = -.139$ , SE = .103,  $\beta^* = .119$ , ns. Approximately 50% of the variance in positive social exchanges,  $R^2 = .436$ , was accounted for by the predictors. The results do not support the possibility of a causal effect in either direction over the thirteen-week period.

## References

Newsom, J.T. (2015). Chapter 13, Missing Data and Attrition in *Longitudinal Structural Equation Modeling: A Comprehensive Introduction*. New York: Routledge.

<sup>&</sup>lt;sup>1</sup> A workaround is to specify single indicator latent variables, each with loading equal to 1 and residual variance equal to 0, and then request the correlation between the latent variables.