#### **APIMoM Output**

Actor-Partner Interdependence Moderation Model Results March 27, 2017

### 1. Text

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#### Summary of APIM Moderation Results

The focus of this study is the investigation of the effect of Other Positivity on Satisfaction being moderated by Tension within the Actor-Partner Interdependence Model or APIM. All three variables are mixed variables and so the relationship between any two variables includes actor and partner effects. The variable names in the dataset are as follows: Other Positivity for is OtherPos\_W, Other Positivity for is OtherPos\_H, Satisfaction for is Satisfaction\_W, Satisfaction for is Satisfaction\_H, Tension for is Tension\_W, and Tension for is Tension\_H. The total number of dyads is 148, and there are no missing data. The dyad members are treated as if they were indistinguishable. The test of distinguishability which includes 8 equality constraints on the coefficients, 5 on the means, 5 on the variances, and 12 on the covariances is statistically significant (chi-square(30) = 44.51, p = .043), with an RMSEA of 0.057. The structural equation models are estimated using the program lavaan. Standardized estimates use pooled variances across members and for interaction effects the standardization uses the product of the two standard deviations and not standard deviation of the product. The X and M variables have been centered, i.e., the mean has been subtracted from all the scores before conducting the moderation analysis. However, the X-axis for figures use the uncentered Other Positivity variable. The descriptive statistics are contained in Table 1.

For the estimates below to be valid, it must be assumed that there is no measurement error in Other Positivity and Tension. Additionally, it must be assumed that there are no unmeasured common causes (i.e., confounders) between Other Positivity and Tension, between Other Positivity and Satisfaction, and between Tension and Satisfaction. It must be assumed that Satisfaction does not cause Other Positivity or Tension and that Tension does not cause Other Positivity. Finally, it must be assumed that the relationships between Other Positivity and Satisfaction and between Tension and Satisfaction are linear and that the interaction between Other Positivity and Satisfaction is also linear.

The combined test the four moderation effects involves fitting two models, one with interaction effects and one without those effects. This combined test of interaction is statistically significant (chi-square(4) = 31.29, p < .001), with an RMSEA of 0.215. Because the RMSEA is greater than .10 and the chi square is statistically significant, there is sufficient evidence to believe that there is moderation.

Table 2 presents the effects in the moderation model. The multiple correlation for the Satisfaction equations is .740. First considered are the "main effects" of Other Positivity on Satisfaction. These are the effects of Other Positivity when the scores of both members on

Tension equal zero. The actor effect equals 0.235 (p < .001) with a standardized effect of .326. The partner effect equals 0.166 (p < .001) with a standardized effect of .230. Next considered are the "main" effects of Tension on Satisfaction. These are the effects of Tension when the scores of both members on Other Positivity equal zero. The actor effect equals -0.277 (p < .001) with a standardized effect of -.278. The partner effect equals -0.098 (p = .002) with a standardized effect of -.099. The variables Other Positivity and Tension interact to affect Satisfaction in four different ways. Each interaction involves two components, the first being Other Positivity and the second being Tension. Thus, the term actor-partner refers to the interaction between the actor variable of Other Positivity and the partner variable of Tension. The actor-actor effect equals 0.185 (p = .002) with a standardized effect of .127. The actor-partner effect equals 0.133 (p = .031) with a standardized effect of .092. The partner-actor effect equals 0.194 (p = .002) with a standardized effect of .133. The partner-partner effect equals 0.032 (p = .591) with a standardized effect of .022.

To better understand these interactions, Table 3 and Figures 1 and 2 present the effects of Other Positivity on Satisfaction for different values of Tension. Simple slopes are computed when either the actor or the partner scores on Tension are either one standard deviation below or one standard deviation above the mean. Because one standard deviation of the moderator equals 0.686, one standard deviation below the mean for Tension is a score of 1.745 and one standard deviation above the mean is a score of 3.117. To examine the actor-actor interaction effect, which is statistically significant (p = .002), the actor effect of Other Positivity on Satisfaction is measured when the actor variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.108 (p = .066) and for one standard deviation above the mean is 0.362 (p < .001). Both of these effects are positive and the Other Positivity actor effect increases as the actor variable for Tension increases. To examine the actor-partner interaction effect, which is statistically significant (p = .031), the actor effect of Other Positivity on Satisfaction is measured when the partner variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.144 (p = .018) and for one standard deviation above the mean is 0.327 (p < .001). Both of these effects are positive and the Other Positivity actor effect increases as the partner variable for Tension increases. To examine the partner-actor interaction effect, which is statistically significant (p = .002), the partner effect of Other Positivity on Satisfaction is measured when the actor variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.033 (p = .588) and for one standard deviation above the mean is  $0.299 \ (p < .001)$ . Both of these effects are positive and the Other Positivity partner effect increases as the actor variable for Tension increases. To examine the partner-partner interaction effect, which is not statistically significant (p = .591), the partner effect of Other Positivity on Satisfaction is measured when the partner variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.144 (p = .014) and for one standard deviation above the mean is 0.188 (p = .001). Both of these effects are positive and the Other Positivity partner effect increases as the partner variable for Tension increases.

To further assist in the understanding of these interactions, Table 4 and Figures 3 and 4 present the effects of Other Positivity on Satisfaction for different actor and partner values of Tension. Simple actor and partner slopes are computed when the actor and partner scores on Tension are both one standard deviation below and one standard deviation above the mean.

Model with A Prior Values for the k's

The user has requested to estimate a model in which constraints are placed on interaction effects to create a simpler and more interpretable model. That constant or k for the effect from interaction Other Positivity on Satisfaction has been set to 1.000 and that constant or k from interaction Tension to Satisfaction has been set to 1.000. The test of the model that imposes these interaction constraints is not statistically significant (chi-square(3) = 5.01, p = .171), with an RMSEA of 0.067. Because the RMSEA is less than .10 and the chi square is not statistically significant, there is evidence that these constraints explain the pattern of interaction effects.

Table 5 presents the effects in the moderation model in which there are constraints on the interaction effects. The multiple correlation for the Satisfaction equations is .736. First considered are the "main" effects of Other Positivity on Satisfaction. These are the effects of Other Positivity when the scores of both members on Tension equal zero. The actor effect equals 0.241 (p < .001) with a standardized effect of .334. The partner effect equals 0.162 (p < .001) with a standardized effect of .224. Next considered are the "main" effects of Tension on Satisfaction. These are the effects of Tension when the scores of both members on Other Positivity equal zero. The actor effect equals -0.281 (p < .001) with a standardized effect of -.282. The partner effect equals -0.091 (p = .003) with a standardized effect of -.092. The variables Other Positivity and Tension interact to affect Satisfaction in four different ways. Each interaction involves two components, the first being Other Positivity and the second being Tension. Thus, the term actor-partner refers to the interaction between the actor variable of Other Positivity and the partner variable of Tension. The actor-actor effect equals 0.136 (p < .001) with a standardized effect of .093. The actor-partner effect equals 0.136 (p < .001) with a standardized effect of .093. The partner-partner effect equals 0.136 (p < .001) with a standardized effect of .093. The partner-partner effect equals 0.136(p < .001) with a standardized effect of .093.

To better understand these interactions, Table 6 and Figure 5 and 6 present the effects of Other Positivity on Satisfaction for different values of Tension with constraints on the interaction effects. Simple slopes are computed when the actor and partner scores on Tension are one standard deviation below and above the mean. Because one standard deviation of the moderator equals 0.686, one standard deviation below the mean for Tension is a score of 1.745 and one standard deviation above the mean is a score of 3.117. To examine the actor-actor interaction effect, which is statistically significant (p < .001), the actor effect of Other Positivity on Satisfaction is measured when the actor variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.148 (p = .001) and for one standard deviation above the mean is 0.334 (p < .001). Both of these effects are positive and the Other Positivity actor effect increases as the actor variable for Tension increases. To examine the actor-partner interaction effect, which is statistically significant (p = .031), the actor effect of Other Positivity on Satisfaction is measured when the partner variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.148 (p = .001) and for one standard deviation above the mean is 0.334 (p < .001). Both of these effects are positive and the Other Positivity actor effect increases as the partner variable for Tension increases. To examine the partner-actor interaction effect, which is statistically significant (p = .002), the partner effect of Other Positivity on Satisfaction is measured when the actor variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.069 (p = .131) and for one standard deviation above the mean is 0.255 (p < .001). Both of these effects are positive and the Other Positivity partner effect increases as the actor variable for Tension increases. To examine the partner-partner interaction effect, which is not statistically significant (p = .591), the partner effect of Other Positivity on Satisfaction is measured when the partner variable for Tension is one standard deviation below and above the mean. The effect for one standard deviation below the mean is 0.069 (p = .131) and for one standard deviation above the mean is 0.255 (p < .001).

Both of these effects are positive and the Other Positivity partner effect increases as the partner variable for Tension increases.

To further assist in the understanding of these interactions, Table 7 and Figures 7 and 8 present the effects of Other Positivity on Satisfaction for different actor and partner values of Tension. Simple actor and partner slopes are computed when the actor and partner scores on Tension are both one standard deviation above and one standard deviation below the mean.

#### 2. Tables

Table 1: Descriptive Statistics

	Variable	Mean	SD	Minimum	Maximum
Other	Positivity	-0.000	0.498	2.600	5.000
Sa	atisfaction	3.605	0.496	1.167	4.000
	Tension	-0.000	0.686	1.167	4.000

Table 2: Effects in the Moderation Model

Cause	Туре	Estimate	р	value	Lower	95% CI	Upper	Standardized
Other Positivity	Actor	0.235		<.001	0.152	to	0.319	0.326
	Partner	0.166		<.001	-0.159	to	0.249	0.230
Tension	Actor	-0.277		<.001	-0.338	to	-0.216	-0.278
	Partner	-0.098		.002	-0.159	to	-0.037	-0.099
Interaction	Actor-Actor	0.185		.002	0.070	to	0.300	0.127
	Actor-Partner	0.133		.031	0.012	to	0.254	0.092
	Partner-Actor	0.194		.002	0.073	to	0.314	0.133
	Partner-Partner	0.032		.591	-0.083	to	0.147	0.022

Table 3: Effects of Other Positivity with Either the Actor or the Partner Effects of Tension +1 (and -1

Type of Effect M for Actor M for Partner Estimate p value Lower 95% CI Upper

Actor	-1sd	mean	0.108	.066 -0.007	to	0.224
	+1sd	mean	0.362	<.001 0.249	to	0.476
	mean	-1sd	0.144	.018 0.024	to	0.264
	mean	+1sd	0.327	<.001 0.212	to	0.441
Partner	-1sd	mean	0.033	.588 -0.087	to	0.153
	+1sd	mean	0.299	<.001 0.184	to	0.414
	mean	-1sd	0.144	.014 0.029	to	0.260
	mean	+1sd	0.188	.001 0.074	to	0.301

Table 4: Effects of Other Positivity with Both Actor and Partner the Effects of Tension +1 (and -1) Sta

Type of Effect	M for Actor M fo	r Partner	Estimate	p value	Lower	95% CI	Upper
Actor	-1sd	-1sd	0.017	.793	-0.112	to	0.147
	-1sd	+1sd	0.200	.002	0.075	to	0.324
	+1sd	-1sd	0.271	<.001	0.144	to	0.398
	+1sd	+1sd	0.454	<.001	0.332	to	0.576
Partner	-1sd	-1sd	0.012	.861	-0.118	to	0.141
	-1sd	+1sd	0.277	<.001	0.149	to	0.406
	+1sd	-1sd	0.055	.402	-0.073	to	0.183
	+1sd	+1sd	0.321	<.001	0.199	to	0.443

Table 4: Effects of Other Positivity with Both Actor and Partner the Effects of Tension +1 (and -1) Star

Cause	Туре	Estimate	p value	Lower	95% CI	Upper	Standardized
Other Positivity	Actor	0.241	<.001	0.158	to	0.324	.334
	Partner	0.162	<.001	-0.152	to	0.245	.224
Tension	Actor	-0.281	<.001	-0.342	to	-0.220	282
	Partner	-0.091	.003	-0.152	to	-0.030	092
Interaction	Actor-Actor	0.136	<.001	0.086	to	0.185	.093
	Actor-Partner	0.136	<.001	0.086	to	0.185	.093
	Partner-Actor	0.136	<.001	0.086	to	0.185	.093
	Partner-Partner	0.136	<.001	0.086	to	0.185	.093

Table 6: Effects of Other Positivity with Either the Actor or the Partner Effects of Tension +1 (and -

Type of Effect N	M for Actor M for	Partner	Estimate	p value	Lower	95% CI	Upper
Actor	-1sd	mean	0.148	.001	0.059	to	0.237
	+1sd	mean	0.334	<.001	0.243	to	0.425
	mean	-1sd	0.148	.001	0.059	to	0.237
	mean	+1sd	0.334	<.001	0.243	to	0.425
Partner	-1sd	mean	0.069	.131	-0.021	to	0.158
	+1sd	mean	0.255	<.001	0.164	to	0.345
	mean	-1sd	0.069	.131	-0.021	to	0.158
	mean	+1sd	0.255	<.001	0.164	to	0.345

Table 7: Effects of Other Positivity with Both the Actor and the Partner Effects of Tension +1 (and -1

Type of Effect	M for Actor M for	Partner	Estimate	p value	Lower	95% CI	Upper
Actor	-1sd	-1sd	0.055	.310	-0.051	to	0.161
	-1sd	+1sd	0.241	<.001	0.134	to	0.348
	+1sd	-1sd	0.241	<.001	0.134	to	0.348
	+1sd	+1sd	0.427	<.001	0.318	to	0.536
Partner	-1sd	-1sd	-0.024	.713	-0.155	to	0.106
	-1sd	+1sd	0.162	.003	0.054	to	0.269
	+1sd	-1sd	0.162	.003	0.054	to	0.269
	+1sd	+1sd	0.348	<.001	0.239	to	0.456

## 3. lavaan Computer Output

Moderation Run with Indistinguishable Dya	ds
lavaan (0.5-22) converged normally after	84 iterations
Number of observations	148
Number of missing patterns	1
Estimator	ML
Minimum Function Test Statistic	44.505
Degrees of freedom	30
P-value (Chi-square)	0.043
lhs op rhs label est se	z pvalue ci.lower ci.upper

1	vv1	~	mv1	ah	-0.277	0 031	-8.891	0.000	-0.338	-0.216
2	yv1 yv2	~	mv1 mv2		-0.277		-8.891	0.000	-0.338	-0.216
3	yv1	~	mv2		-0.098		-3.156	0.002	-0.159	-0.037
4	yv2	~	mv1	-	-0.098		-3.156	0.002	-0.159	-0.037
5	yv1	~	xv1	aa		0.042	5.556	0.000	0.152	0.319
6	yv2	~	xv2	aa		0.042	5.556	0.000	0.152	0.319
7	yv1	~	xv2	pa		0.042	3.919	0.000	0.083	0.249
8	yv2	~	xv1	pa		0.042	3.919	0.000	0.083	0.249
9	yv1	~	xm11	iAA		0.059	3.154	0.002	0.070	0.300
10	yv2	~	xm11	iPP		0.059	0.537	0.591	-0.083	0.147
11	yv1	~	xm12	iAP		0.062	2.158	0.031	0.012	0.254
12	yv2	~	xm12	iPA		0.062	3.144	0.002	0.073	0.314
13	yv1	~	xm21	iPA		0.062	3.144	0.002	0.073	0.314
14	yv2	~	xm21	iAP		0.062	2.158	0.031	0.012	0.254
15	yv1	~	xm22	iPP		0.059	0.537	0.591	-0.083	0.147
16	yv2	~	xm22	iAA	0.185	0.059	3.154	0.002	0.070	0.300
17	xv1 -	~ ~	xv2		0.057	0.021	2.744	0.006	0.016	0.098
18	yv1 -	~ ~	yv2		0.033	0.010	3.470	0.001	0.014	0.052
19	mv1 ~	~ ~	mv2		0.149	0.041	3.670	0.000	0.069	0.228
20	$xm11$ $\cdot$	~ ~	xm22		0.018	0.010	1.703	0.089	-0.003	0.038
21	xm12 ~	~ ~	xm21		0.025	0.009	2.672	0.008	0.007	0.043
22	xv1 ~	~1		m1	0.000	0.032	0.000	1.000	-0.063	0.063
23	xv2 ~	~1		m1	0.000	0.032	0.000	1.000	-0.063	0.063
24	yv1 -	~1		m2	3.662	0.025	147.757	0.000	3.614	3.711
25	yv2 -	~1		m2			147.757	0.000	3.614	3.711
26	mv1 ~	~1		mЗ	0.000	0.046	0.000	1.000	-0.090	0.090
27	mv2 ~			mЗ		0.046	0.000	1.000	-0.090	0.090
	xm11 ~				-0.126		-5.716	0.000	-0.169	-0.083
	xm22 ~				-0.126		-5.716	0.000	-0.169	-0.083
	xm12 ~				-0.093		-4.322	0.000	-0.135	-0.051
	xm21 -				-0.093		-4.322	0.000	-0.135	-0.051
32	xv1 -		xv1	v1		0.021	11.852	0.000	0.207	0.289
33	xv2 ~		xv2	v1		0.021	11.852	0.000	0.207	0.289
34	yv1 -		yv1	v2		0.010	11.660	0.000	0.092	0.130
35	yv2 -		yv2	v2		0.010	11.660	0.000	0.092	0.130
36 37	mv1 ~ mv2 ~		mv1 mv2	v3 v3		0.041	11.599 11.599	0.000	0.391 0.391	0.550 0.550
	xm11 ~		xm11	v3 v4		0.041	12.046	0.000	0.391	0.330
	xm22 ~		xm22	v4 v4		0.010	12.040 12.046	0.000	0.105	0.140 0.146
	xm12 ·		xm12	v-1 v5		0.009	11.869	0.000	0.093	0.140
	xm21 ~		xm21	v5		0.009	11.869	0.000	0.093	0.129
42	xv1 -		xm11	c1		0.010	1.662	0.097	-0.003	0.038
43	xv2 ~		xm22	c1		0.010	1.662	0.097	-0.003	0.038
44	xv2 ~		xm11		-0.004		-0.388	0.698	-0.025	0.016
45	xv1 -		xm22		-0.004		-0.388	0.698	-0.025	0.016
46	xv1 -		xm12	c3	0.012		1.262	0.207	-0.007	0.032
47	xv2 ~		xm21	c3		0.010	1.262	0.207	-0.007	0.032
48	xv2 ~	~ ~	xm12	c4	-0.004	0.010	-0.409	0.682	-0.023	0.015
49	xv1 -	~ ~	xm21	c4	-0.004	0.010	-0.409	0.682	-0.023	0.015
50	mv1 ~	~ ~	xm11	c5	-0.042	0.015	-2.844	0.004	-0.070	-0.013
51	mv2 ~	~ ~	xm22		-0.042		-2.844	0.004	-0.070	-0.013
52	mv2 ~		xm11		-0.016		-1.086	0.277	-0.045	0.013
53	mv1 ~		xm22		-0.016		-1.086	0.277	-0.045	0.013
54	mv1 -	~ ~	xm12	c7	-0.016	0.014	-1.154	0.248	-0.043	0.011

55 mv2 ~~	xm21 c	7 -0.016	0.014	-1.154	0.248	-0.043	0.011
56 mv2 ~~	xm12 c	8 -0.015	0.014	-1.074	0.283	-0.042	0.012
57 mv1 ~~	xm21 c	8 -0.015	0.014	-1.074	0.283	-0.042	0.012
58 xm11 ~~			0.007	5.273		0.025	0.054
59 xm21 ~~			0.007	5.273	0.000	0.025	0.054
60 xm11 ~~			0.007	2.581	0.010	0.005	0.034
61 xm12 ~~			0.007	2.581	0.010	0.005	0.034
62 mv1 ~~		1 -0.126		-5.593	0.000	-0.170	-0.082
63 mv2 ~~		1 -0.126		-5.593		-0.170	-0.082
64 mv2 ~~	xv1 c1	2 -0.093	0.022	-4.119	0.000	-0.137	-0.049
65 mv1 ~~	xv2 c1	2 -0.093	0.022	-4.119	0.000	-0.137	-0.049
66 kx :=	pa/aa k	x 0.705	0.199	3.536	0.000	0.314	1.096
	pb/ab k					0.130	0.579
	std.all						
1 -0.277	-0.383						
2 -0.277	-0.383						
3 -0.098	-0.136						
4 -0.098	-0.136						
5 0.235	0.236						
6 0.235	0.236						
7 0.166	0.167						
8 0.166	0.167						
9 0.185	0.132						
10 0.032	0.023						
11 0.133	0.089						
12 0.194	0.130						
13 0.194	0.130						
14 0.133	0.089						
15 0.032	0.023						
16 0.185	0.132						
17 0.057	0.232						
18 0.033	0.298						
19 0.149	0.316						
20 0.018	0.141						
21 0.025	0.225						
22 0.000	0.000						
23 0.000	0.000						
24 3.662	7.390						
25 3.662	7.390						
26 0.000	0.000						
27 0.000	0.000						
28 -0.126	-0.355						
29 -0.126	-0.355						
30 -0.093	-0.278						
31 -0.093	-0.278						
32 0.248	1.000						
33 0.248	1.000						
34 0.111	0.452						
35 0.111	0.452						
36 0.471	1.000						
37 0.471	1.000						
38 0.125	1.000						
39 0.125	1.000						
40 0.111	1.000						

41 0.111	1 000	ſ							
42 0.017									
43 0.017									
44 -0.004									
45 -0.004	1 -0.023	3							
46 0.012	2 0.07	5							
47 0.012	2 0.07	5							
48 -0.004	1 -0.024	4							
49 -0.004	1 -0.024	4							
50 -0.042	2 -0.172	2							
51 -0.042	2 -0.172	2							
52 -0.016	5 -0.066	6							
53 -0.016	6 -0.066	6							
54 -0.016	5 -0.070	C							
55 -0.016	5 -0.070	C							
56 -0.015	5 -0.06	5							
57 -0.015									
58 0.039									
59 0.039									
60 0.019									
61 0.019									
62 -0.126									
63 -0.126									
64 -0.093									
65 -0.093 66 0.705									
67 0.355									
01 0.000	0.000	5							
Moderatio							_		
lavaan ((	).5-22) (	conver	ged not	rmally	after 7	'2 itera	tions		
Number	of obse	rvatio	ns				148		
Wallber	01 00000		,110				110		
Number	of miss:	ing pa	atterns				1		
Estimat	or						ML		
	n Functio	n Tos	t Stati	istic		49	.520		
	s of free					-10	33		
	e (Chi-so					0	.032		
lhs c		label	est	se	Z			ci.upper st	d.lv
1 yv1	~ mv1		-0.281		-9.005	0.000	-0.342	-0.220 -0	
2 yv2	~ mv2		-0.281		-9.005	0.000	-0.342	-0.220 -0	
3 yv1	~ mv2		-0.091		-2.928	0.003	-0.152	-0.030 -0	
4 yv2	~ mv1	-	-0.091		-2.928	0.003	-0.152	-0.030 -0	0.091
5 yv1	~ xv1	aa		0.042	5.682	0.000	0.158	0.324 0	0.241
6 yv2	~ xv2	aa	0.241	0.042	5.682	0.000	0.158	0.324 0	0.241
7 yv1	~ xv2	pa	0.162	0.042	3.808	0.000	0.078	0.245 0	0.162
8 yv2	~ xv1	pa	0.162	0.042	3.808	0.000	0.078	0.245 0	0.162
			0 126	0.025	5.362	0.000	0.086	0.185 0	).136
9 yv1	~ xm11	iAA	0.136	0.025	0.002	0.000			
9 yv1 10 yv2	~ xm11 ~ xm11	iAA iPP		0.025	5.362	0.000	0.086		0.136
-			0.136					0.185 0	
10 yv2 11 yv1 12 yv2	~ xm11 ~ xm12 ~ xm12	iPP iAP iPA	0.136 0.136	0.025	5.362 5.362 5.362	0.000	0.086	0.185 ( 0.185 (	0.136
10 yv2 11 yv1 12 yv2 13 yv1	~ xm11 ~ xm12 ~ xm12 ~ xm21	iPP iAP iPA iPA	0.136 0.136 0.136	0.025 0.025	5.362 5.362 5.362 5.362	0.000 0.000	0.086 0.086	0.185 ( 0.185 ( 0.185 (	).136 ).136
10 yv2 11 yv1 12 yv2	~ xm11 ~ xm12 ~ xm12	iPP iAP iPA	0.136 0.136 0.136 0.136	0.025 0.025 0.025	5.362 5.362 5.362	0.000 0.000 0.000	0.086 0.086 0.086	0.185 ( 0.185 ( 0.185 ( 0.185 (	).136 ).136 ).136

15 yv1 ~ :		0.136	0.025	5.362	0.000	0.086	0.185 0.136
16 yv2 ~ :	xm22 iAA	0.136	0.025	5.362	0.000	0.086	0.185 0.136
17 xv1 ~~	xv2		0.021	2.744	0.006	0.016	0.098 0.057
18 yv1 ~~	yv2		0.010	3.336	0.001	0.013	0.051 0.032
19 mv1 ~~	mv2	0.149		3.670	0.000	0.069	0.228 0.149
20 xm11 ~~ :	xm22	0.018		1.703	0.089	-0.003	0.038 0.018
21 xm12 ~~ :	xm21	0.025	0.009	2.672	0.008	0.007	0.043 0.025
22 xv1 ~1	m1	0.000	0.032	0.000	1.000	-0.063	0.063 0.000
23 xv2 ~1	m1	0.000	0.032	0.000	1.000	-0.063	0.063 0.000
24 yv1 ~1	m2	3.664	0.025	148.290	0.000	3.615	3.712 3.664
25 yv2 ~1	m2	3.664	0.025	148.290	0.000	3.615	3.712 3.664
26 mv1 ~1	m3	0.000	0.046	0.000	1.000	-0.090	0.090 0.000
27 mv2 ~1	m3	0.000	0.046	0.000	1.000	-0.090	0.090 0.000
28 xm11 ~1	m4	-0.126	0.022	-5.716	0.000	-0.169	-0.083 -0.126
29 xm22 ~1	m4	-0.126	0.022	-5.716	0.000	-0.169	-0.083 -0.126
30 xm12 ~1	m5	-0.093	0.021	-4.322	0.000	-0.135	-0.051 -0.093
31 xm21 ~1	m5	-0.093	0.021	-4.322	0.000	-0.135	-0.051 -0.093
32 xv1 ~~	xv1 v1	0.248	0.021	11.852	0.000	0.207	0.289 0.248
33 xv2 ~~	xv2 v1	0.248	0.021	11.852	0.000	0.207	0.289 0.248
34 yv1 ~~	yv1 v2	0.113	0.010	11.699	0.000	0.094	0.131 0.113
35 yv2 ~~	yv2 v2	0.113	0.010	11.699	0.000	0.094	0.131 0.113
36 mv1 ~~	mv1 v3	0.471	0.041	11.599	0.000	0.391	0.550 0.471
37 mv2 ~~	mv2 v3	0.471	0.041	11.599	0.000	0.391	0.550 0.471
38 xm11 ~~ :	xm11 v4	0.125	0.010	12.046	0.000	0.105	0.146 0.125
39 xm22 ~~ :				12.046	0.000	0.105	0.146 0.125
40 xm12 ~~				11.869	0.000	0.093	0.129 0.111
41 xm21 ~~ :				11.869	0.000	0.093	0.129 0.111
42 xv1 ~~ :				1.662	0.097	-0.003	0.038 0.017
43 xv2 ~~ :				1.662	0.097	-0.003	0.038 0.017
44 xv2 ~~ :		-0.004		-0.388	0.698	-0.025	0.016 -0.004
45 xv1 ~~		-0.004		-0.388	0.698	-0.025	0.016 -0.004
46 xv1 ~~				1.262	0.207	-0.007	0.032 0.012
47 xv2 ~~				1.262	0.207	-0.007	0.032 0.012
48 xv2 ~~		-0.004		-0.409	0.682	-0.023	0.015 -0.004
49 xv1 ~~		-0.004		-0.409	0.682	-0.023	0.015 -0.004
50 mv1 ~~		-0.042		-2.844	0.004	-0.070	-0.013 -0.042
51 mv2 ~~		-0.042		-2.844	0.004	-0.070	-0.013 -0.042
52 mv2 ~~		-0.016		-1.086	0.277	-0.045	0.013 -0.016
53 mv1 ~~		-0.016		-1.086	0.277	-0.045	0.013 -0.016
54 mv1 ~~		-0.016		-1.154	0.248	-0.043	0.011 -0.016
55 mv2 ~~ :		-0.016		-1.154	0.248	-0.043	0.011 -0.016
56 mv2 ~~ :		-0.015		-1.074	0.283	-0.042	0.012 -0.015
57 mv1 ~~ :		-0.015		-1.074	0.283	-0.042	0.012 -0.015
58 xm11 ~~ :				5.273	0.000	0.012	0.054 0.039
59 xm21 ~~				5.273	0.000	0.025	0.054 0.039
60 xm11 ~~				2.581	0.010	0.025	0.034 0.019
61 xm12 ~~ :				2.581	0.010	0.005	0.034 0.019
62 mv1 ~~		-0.126		-5.593	0.010	-0.170	-0.082 -0.126
63 mv2 ~~		-0.126		-5.593	0.000	-0.170	-0.082 -0.126
64 mv2 ~~		-0.093		-4.119	0.000	-0.137	-0.049 -0.093
65 mv1 ~~		-0.093			0.000	-0.137	-0.049 - 0.093
std.all	AVZ CIZ	0.033	0.022	7.113	0.000	0.137	0.040 0.000
1 -0.389							

1 -0.389

2 -0.389

$\begin{array}{c} 3\\ 4\\ 5\\ 6\\ 7\\ 8\\ 9\\ 10\\ 11\\ 12\\ 13\\ 14\\ 15\\ 16\\ 17\\ 18\\ 19\\ 20\\ 12\\ 23\\ 24\\ 25\\ 26\\ 27\\ 28\\ 29\\ 30\\ 31\\ 23\\ 34\\ 35\\ 6\\ 37\\ 38\\ 9\\ 40\\ 41\\ 24\\ 34\\ 45\\ 46\\ 47\\ 48\\ 9\\ 50\\ 10\\ 10\\ 10\\ 10\\ 10\\ 10\\ 10\\ 10\\ 10\\ 1$	-0.126 -0.126 0.242 0.242 0.162 0.097 0.097 0.091 0.091 0.091 0.091 0.097 0.232 0.285 0.316 0.141 0.225 0.300 0.000 7.393 7.393 0.000 0.000 7.393 7.393 0.000 0.000 7.393 7.393 0.000 0.000 7.393 7.393 0.000 0.000 7.393 7.393 0.000 0.000 0.000 7.393 7.393 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000
47 48 49	0.075 -0.024 -0.024 -0.172 -0.172 -0.066 -0.066 -0.070 -0.070
56	-0.065

57	-0.065
58	0.331
59	0.331
60	0.162
61	0.162
62	-0.368
63	-0.368
64	-0.271
65	-0.271

# 4. Figures

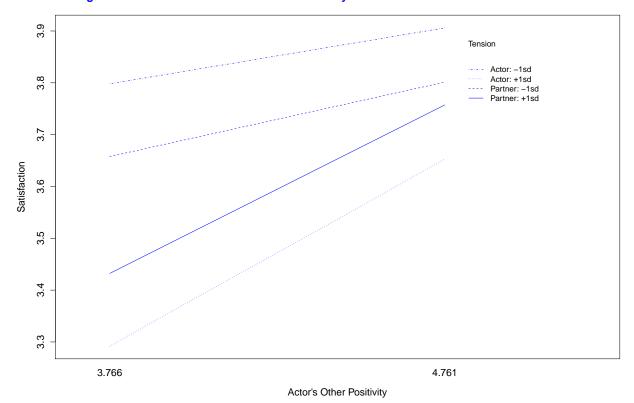
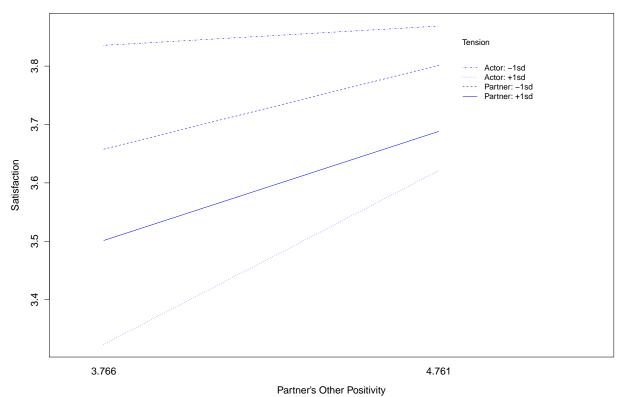


Figure 1: The Effect for the Actor's Other Positivity at Different Actor or Partner Moderator Values





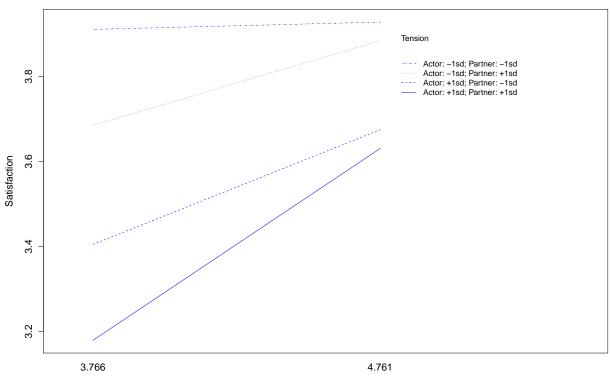


Figure 3: The Effect for the Actor's Other Positivity at Different Actor and Partner Moderator Values

Actor's Other Positivity

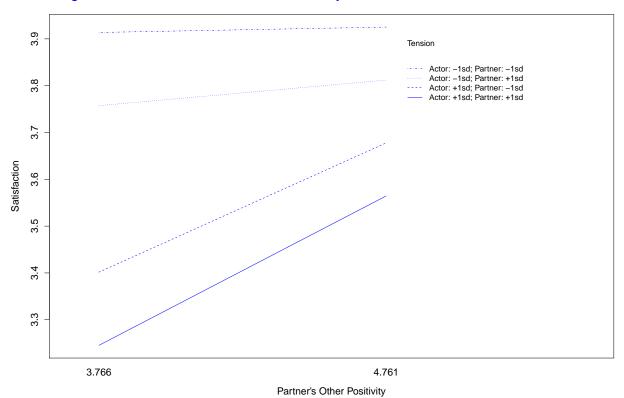


Figure 4: The Effect for the Partner's Other Positivity at Different Actor and Partner Moderator Values

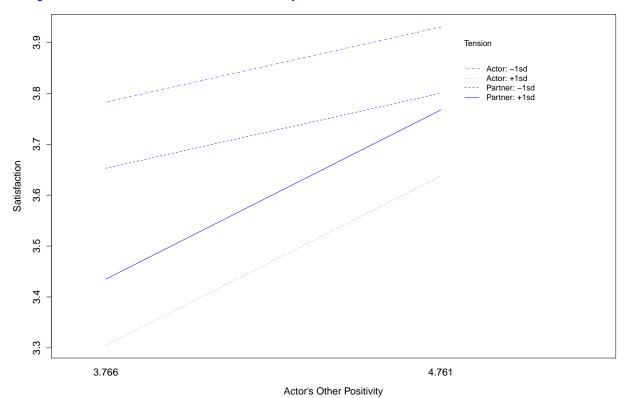


Figure 5: The Effect for the Actor's Other Positivity with Constraints at Different Actor or Partner Moderator Value:

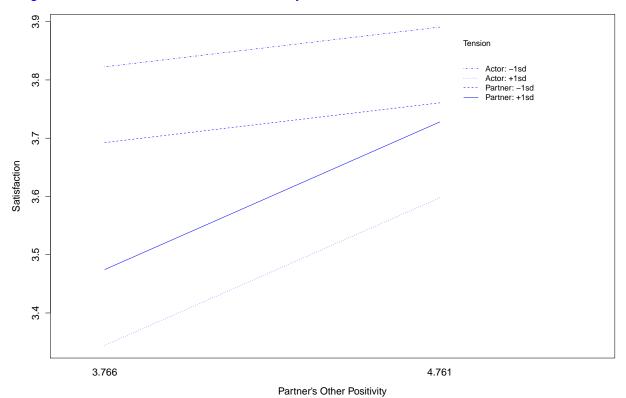


Figure 6: The Effect for the Partner's Other Positivity with Constraints at Different Actor or Partner Moderator Value

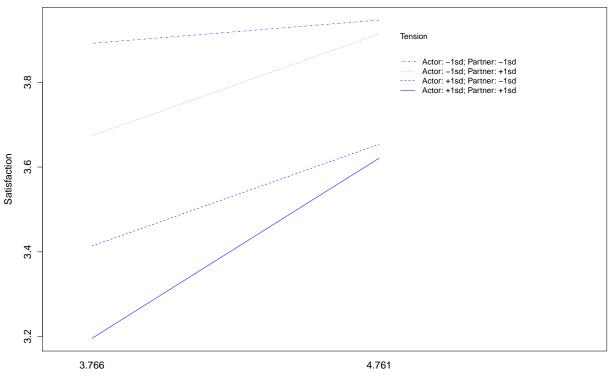
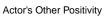


Figure 7: The Effect for the Actor's Other Positivity with Constraints at Different Actor and Partner Moderator Value



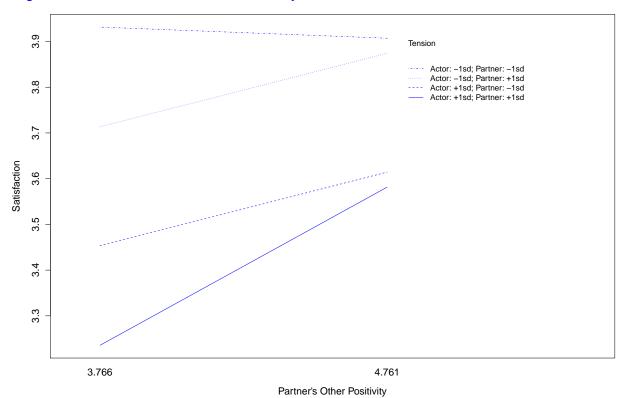


Figure 8: The Effect for the Partner's Other Positivity with Constraints at Different Actor and Partner Moderator Valu