Correlation Matrix

		SBS_Total	Positive Affect	Negative Affect
SBS_Total	**Spearman's rho**	—		
	df	—		
	p-value	—		
Positive Affect	**Spearman's rho**	-0.092	—	
	df	207	—	
	p-value	0.187	—	
Negative Affect	**Spearman's rho**	0.328***	-0.272***	—
	df	207	207	—
	p-value	<.001	<.001	—

Note. * p < .05, ** p < .01, *** p < .001