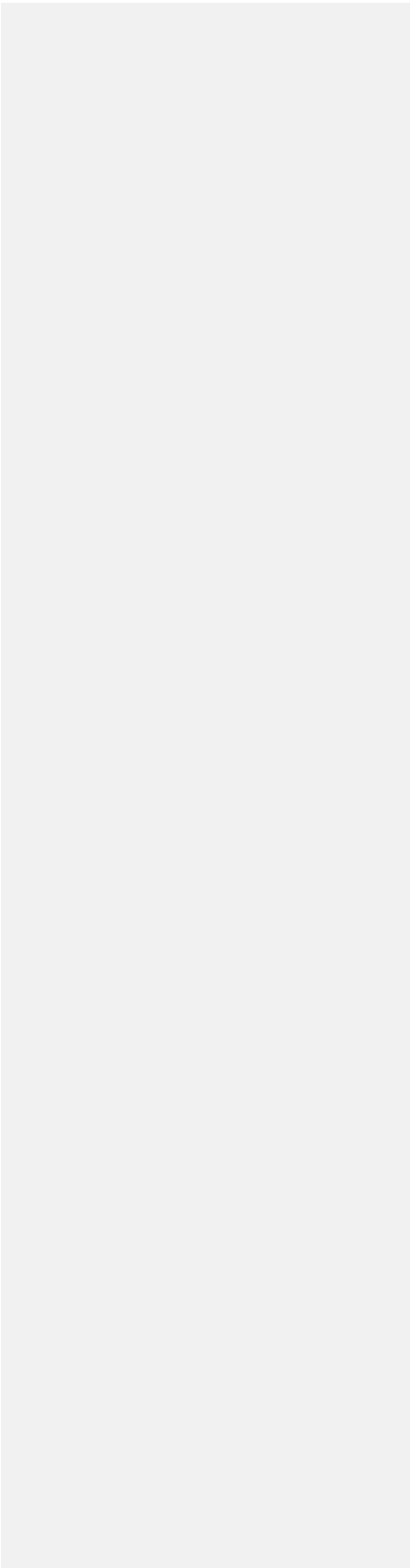


PROPOSED SUPPLEMENTARY ONLINE TABLES (2)



Supplementary Table 1

Confirmatory factor analysis goodness-of-fit indices for the MEIM-R by modality in the GEM trial analytic sample

Model	$\chi^2_{SB} (df)$	SRMR	RMSEA [90% CI]	CFI	AIC	CFI difference	AIC difference	$\Delta\chi^2_{SB} (df)$
<i>Measurement invariance across modalities</i>								
1. Fit of correlated two-factor model								
1a. Mail ($n = 741$)	32.61 (8)***	.03	.06 [.04-.09]	.98	10311.33			
1b. Online ($n = 722$)	43.06 (8)***	.02	.08 [.06-.10]	.98	10073.74			
2. Equal factor structure	75.81 (16)***	.03	.07 [.06-.09]	.98	20385.08			
3. Equal factor loadings	80.28 (20)***	.03	.06 [.05-.08]	.98	20378.95	.000	-6.13	1.59 (4)
4. Equal item intercepts	88.05 (16)***	.03	.06 [.05-.07]	.98	20375.81	.001	-3.14	4.90 (4)

Note. χ^2_{SB} = Satorra–Bentler scaled chi-square; SRMR = standardized root mean square residual; RMSEA = root mean square error of approximation; CI = confidence interval; CFI = Bentler comparative fit index; AIC = Akaike Information Criterion.

*** $p < .001$.

Supplementary Table 2

Confirmatory factor analysis parameter estimates, means, standard deviations, and Cronbach's alphas for the MEIM-R by modality in the GEM trial analytic sample

		CFA completely standardized parameter estimates ^a													
Group	n	Exploration				Commitment			Exploration subscale		Commitment subscale		Overall scale		
		A	E1	E4	E5	C2	C3	C6	M (SD)	α	M (SD)	α	M (SD)	α	
Mail	741	.72	.56	.91	.79	.84	.81	.88	3.17 (0.90)	.79	3.76 (0.85)	.88	3.47 (0.77)	.86	
Online	722	.78	.64	.94	.83	.89	.85	.89	3.09 (1.00)	.86	3.60 (0.96)	.91	3.34 (0.89)	.90	

Note. Means and standard deviations represent observed scores.

^aLabels correspond to paths in Figure 1.