

Supplemental Information

Table S7. Standardized total, direct and indirect effects for genetic ancestry and self-reported race in test null and alternate models.

Model	Genetic Ancestry				Self-Reported Race			
	Direct		Indirect		Direct		Indirect	
	B (s.e. B)	β	B	β	B (s.e. B)	β	B	β
Model 1 - full	NA		NA		-.037 (.016)	-.050*	-.031	-.043*
Model 2 - full	-.388(.177)	-.066*	-.195	-.033	-.004 (.022)	-.005	-.030	-.041
Model 3- reduced	-.387 (.177)	-.066*	-.194	-.033	-.005 (.022)	-.006	-.029	-.040
Model 4 - final	-.581 (.194)	-.100**	-.167	-.029	-.012 (.024)	-.017	NA	NA
Model 4 (HRSD ₁₇)	-.604 (.190)	-.108***	-.142	-.025	-.010 (.023)	-.014	NA	NA

Note. Standardized Coefficients (β) are directly comparable within models across predictors, and represent the standard deviation change in outcome for one standard deviation change in the predictors. Unstandardized coefficients (B) represent the unit change in the criterion (Treatment Response) raw score for each unit change in the predictor raw score and are not directly comparable across predictors.

* $p \leq .05$; ** $p \leq .01$, *** $p \leq .001$