

Supplementary Online Content

Greenwood TA, Lazzeroni LC, Maihofer AX, et al. Genome-wide association of endophenotypes for schizophrenia from the Consortium on the Genetics of Schizophrenia (COGS) study. *JAMA Psychiatry*. Published online October 9, 2019. doi:10.1001/jamapsychiatry.2019.2850

eTable. Correlations Among All 11 Endophenotypes

eFigure. Quantile-Quantile Plots for Each of the 11 Endophenotypes

This supplementary material has been provided by the authors to give readers additional information about their work.

eTable. Correlations Among All 11 Endophenotypes

	PPI	AS	CVLT	DS-CPT	LNS	ABF	FMEM	SMEM	SPA	S-M
AS	0.13 ^c									
CVLT	0.05	0.45^c								
DS-CPT	0.13 ^c	0.33^c	0.31^c							
LNS	0.10 ^a	0.41^c	0.52^c	0.27 ^c						
ABF	0.14 ^c	0.32^c	0.35^c	0.22 ^c	0.37^c					
FMEM	0.05	0.35^c	0.35^c	0.21 ^c	0.32^c	0.36^c				
SMEM	0.09 ^a	0.34^c	0.34^c	0.20 ^c	0.33^c	0.46^c	0.57^c			
SPA	0.08 ^a	0.43^c	0.40^c	0.28 ^c	0.43^c	0.49^c	0.46^c	0.44^c		
S-M	0.05	0.42^c	0.39^c	0.22 ^c	0.36^c	0.43^c	0.50^c	0.44^c	0.56^c	
EMO	0.12 ^b	0.45^c	0.42^c	0.24 ^c	0.42^c	0.43^c	0.57^c	0.47^c	0.54^c	0.59^c

All endophenotype values used for comparison represent standardized residuals adjusted for age (AS, CVLT, DS-CPT, LNS, ABF, FMEM, SMEM, SPA, S-M, EMO) and/or sex (PPI, CVLT, ABF, FMEM, EMO). Significance is indicated as $p < 0.05^a$, $p < 0.01^b$, or $p < 0.005^c$, the latter of which reflects a conservative Bonferroni correction for multiple comparisons of independent variables. Moderate correlations ($r > 0.3$) are shown in bold text.

eFigure. Quantile-Quantile Plots for Each of the 11 Endophenotypes

