

Supplementary Figure 6a-c. Correlations between 13 genome-wide polygenic scores. These results are based on GPS constructed using a GWAS $P_T = 0.30, 0.10,$ and 0.05 . P -values that pass Bonferroni correction are indicated with two asterisks, while those reaching nominal significance (thus suggestive evidence) are shown with a single asterisk.