

Figure S6. PheWAS meta-analysis results for the trans-ethnic TC PGS in UK Biobank and MVP. The blue horizontal line denotes phenome-wide significance ($p <= 6.5 \times 10^{-05}$) and the red line is genome-wide significance ($p <= 5 \times 10^{-08}$). Phenotypes have been pruned, so that the most significant one per correlated phenotype group (correlation coefficient > 0.2) is retained. Pairwise correlations were estimated with chi-square test and Cramer's V for the dichotomous phenotypes and Pearson's correlation for the continuous phenotypes. Full phenome-wide significant results are presented in Additional file 12: Table S8.