

Supplementary Table 3. Results of GCTA

	β_0 (intercept)	β_1 (slope)
Genetic variance (SE)	5.34 (1.97)	0.24 (0.17)
Residual variance (SE)	50.94 (2.06)	4.89 (0.18)
Phenotypic variance (SE)	56.29 (0.77)	5.14 (0.07)
Ratio of genetic variance to phenotypic variance (SE)	0.10 (0.03)	0.04 (0.03)
<i>P</i> value	3.52×10^{-3}	7.58×10^{-2}

GCTA, genome-wide complex trait analysis; SE, standard error.