Table S14. Genomewide h_g^2 and h_{gLD}^2 of liability for all case-control traits.

Phenotype	Genotyped		Genotyped & imputed	
	h_g^2 total (se)	h_{gLD}^2 total (se)	h_g^2 total (se)	h_{gLD}^2 total (se)
BD	0.26 (0.032)	0.27 (0.047)	0.19 (0.028)	0.27 (0.063)
CAD	0.30 (0.058)	0.31 (0.085)	0.20 (0.048)	0.32 (0.112)
CD	0.18 (0.024)	0.20 (0.036)	0.10 (0.021)	0.25 (0.047)
HT	0.60 (0.089)	0.82 (0.131)	0.36 (0.079)	0.88 (0.172)
RA	0.11 (0.031)	0.17 (0.047)	0.04 (0.026)	0.15 (0.061)
T1D	0.13 (0.030)	0.16 (0.045)	0.08 (0.026)	0.14 (0.059)
T2D	0.36 (0.066)	0.55 (0.097)	0.28 (0.055)	0.59 (0.127)
UC	0.17 (0.017)	0.25 (0.028)	-	-
MS	0.19 (0.009)	0.26 (0.014)	-	-