

Supplemental Figure S8. Global QQ plots for association analysis of rare variants with LDL in the UK10K TWINSUK cohort: (a) FunSPU (genomic control $\lambda = 0.611$), (b) FunSPUw ($\lambda = 0.805$), (c) wtFunSPU with global weights ($\lambda = 0.642$), and (d) wtFunSPUw with global weights ($\lambda = 0.789$).

