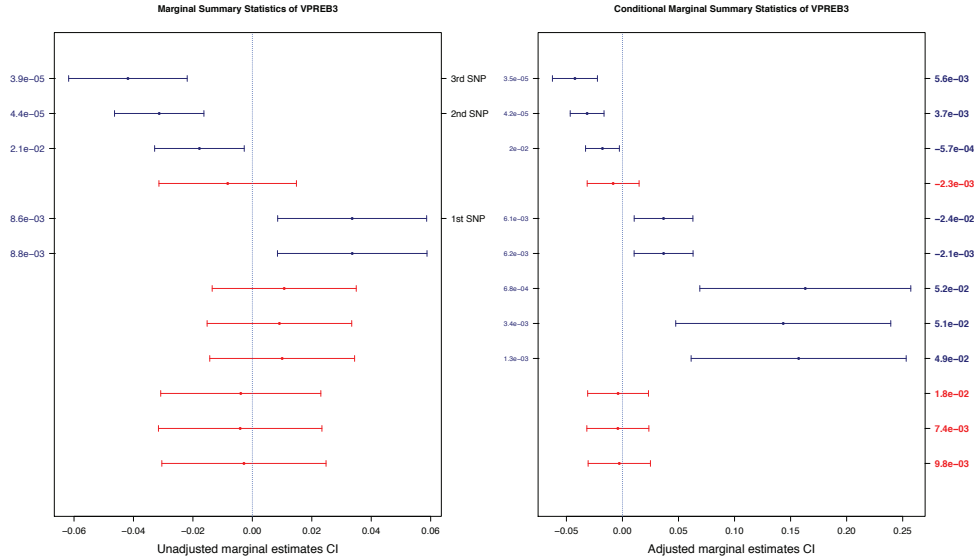

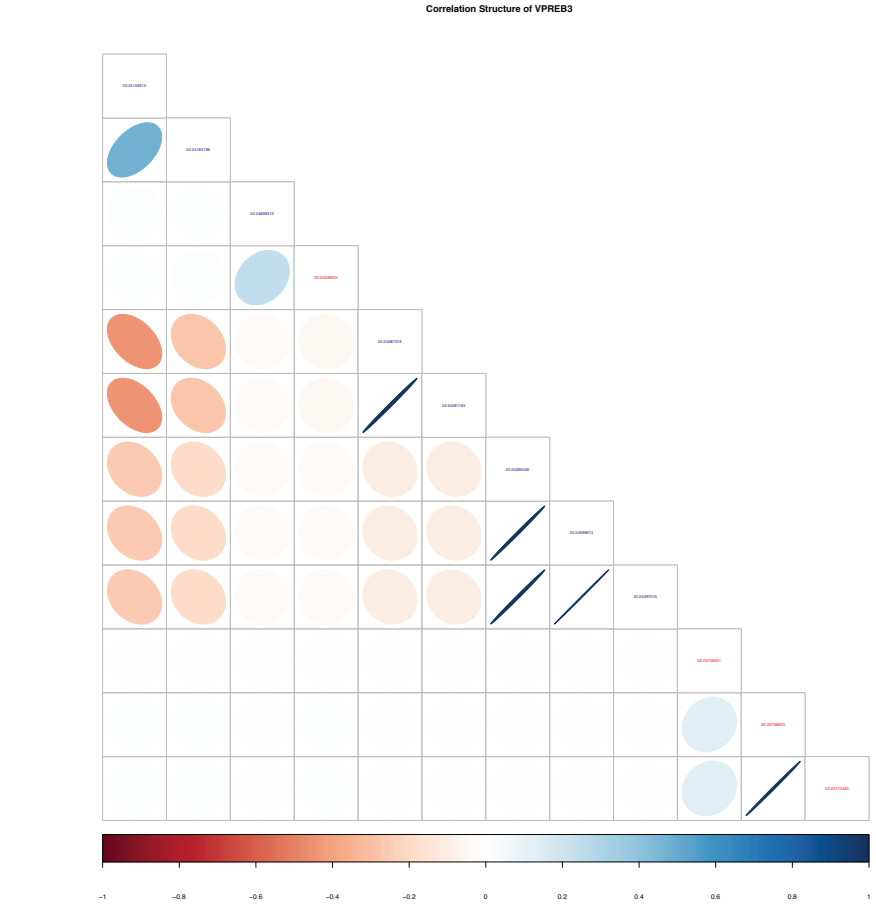

Fig S2. Gene *VPREB3*: (a) forest plot of marginal association of genetic variants in *VPREB3*; (b) forest plot of conditional association adjusting for predicted gene expression; (c) pairwise linkage disequilibrium (LD) R^2. The p-values < 0.05 are labeled on the left margin in (a) and (b). The weights used in calculating gene expression are labeled on the right margin in (b).