



On the X -axis genomic positions of SNPs aligned to NCBI genome build 36, chromosome borders are designated by changing dot colors. On the Y -axis $-\log_{10}(p \text{ values})$ for association between SNP genotype and disease status as obtained from logistic regression analyses in the GWAS discovery set. The dotted line indicates the threshold for genome-wide significance ($p=5 \times 10^{-8}$). GWAS, genome-wide association study.