**Supplementary Fig. 6.** Search result of rs10965250 in Korean Reference Genome Database (KRG-DB). On the variant, there is a genome-wide significant result for type 2 diabetes mellitus development in the genome-wide association study (GWAS) catalog. In the genome-wide repository of associations between SNPs and phenotypes (GRASP), many results having nominal significance ($P<0.05$) are shown. NCBI, National Center for Biotechnology Information.