



Supplementary Fig. 2. Genome-wide association study (GWAS) results of rs2237892 from Korean Reference Genome Database (KRGDB). Search result from KRGDB. The original data comes from GWAS catalog and genome-wide repository of associations between SNPs and phenotypes (GRASP) database. The results from GWAS catalog had genome-wide significance ($P < 5 \times 10^{-8}$), while those from GRASP had a range of P value from nominal significance to genome-wide significance.