Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_PD_CASE_CONTROL_PARENT_PROXY_2021

LOF (LOFTEE) and CADD > 20 (VEP) GBA CMCWald Burden Test with MAF \leq 0.01 and N_{variants per gene} \geq 3 20 15 $-\log_{10}(p)$ 10 5 2 3 10 12 13 14 15 17 18 19 20 2122

Chromosome