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

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_PD_CASE_CONTROL_PROXIES_2021

LOF (LOFTEE)

CMCWald Burden Test with MAF $\leq$ 0.01 and $N_{variants\ per\ gene} \geq 1$