

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

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_ALL_PD_PHENOTYPES_CONTROL_2021

GBA

Moderate or High Impact (SnpEff)

CMCWald Burden Test with $MAF \leq 0.01$ and $N_{\text{variants per gene}} \geq 2$

