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

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_PD_CASE_CONTROL_2021

Missense (SnpEff)

CMCWald Burden Test with MAF ≤ 0.001 and N$_{\text{variants per gene}} \geq 3$