

Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES
meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_PD_CASE_CONTROL_PROXIES_2021

Moderate or High Impact (Snpeff)

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

