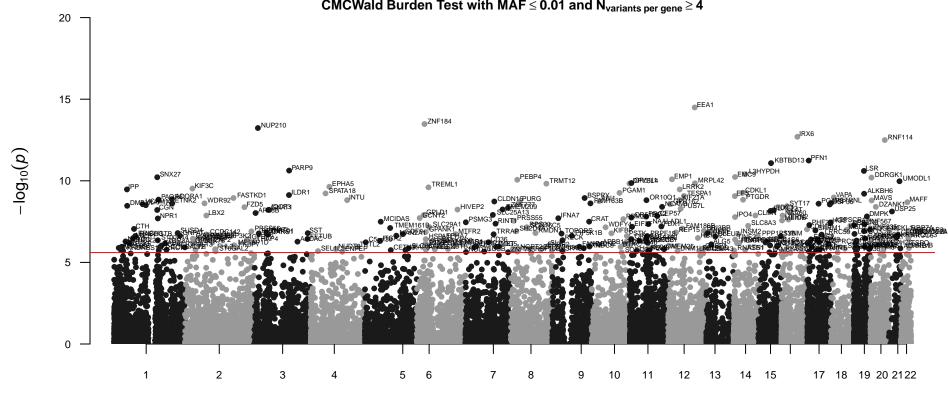
Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_PD_CASE_CONTROL_PARENT_PROXY_2021

Moderate or High Impact (SnpEff)

CMCWald Burden Test with MAF ≤ 0.01 and N_{variants per gene} ≥ 4



Chromosome