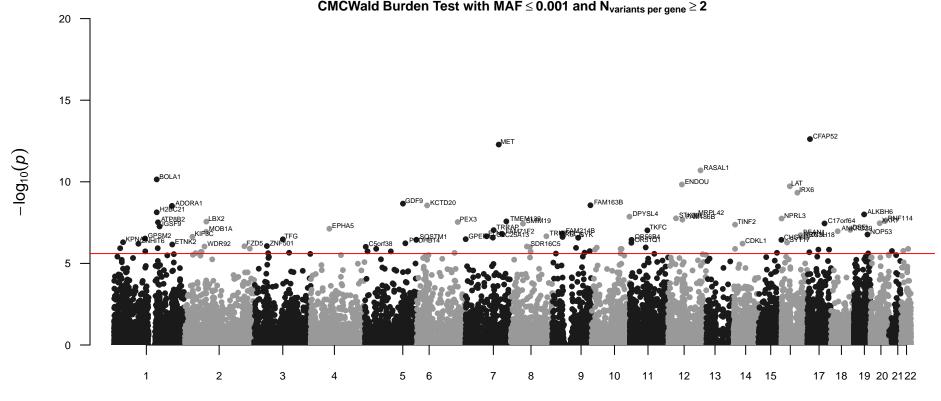
Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_ALL_PD_PHENOTYPES_CONTROL_2021

Moderate or High Impact (SnpEff)

CMCWald Burden Test with MAF ≤ 0.001 and N_{variants per gene} ≥ 2



Chromosome