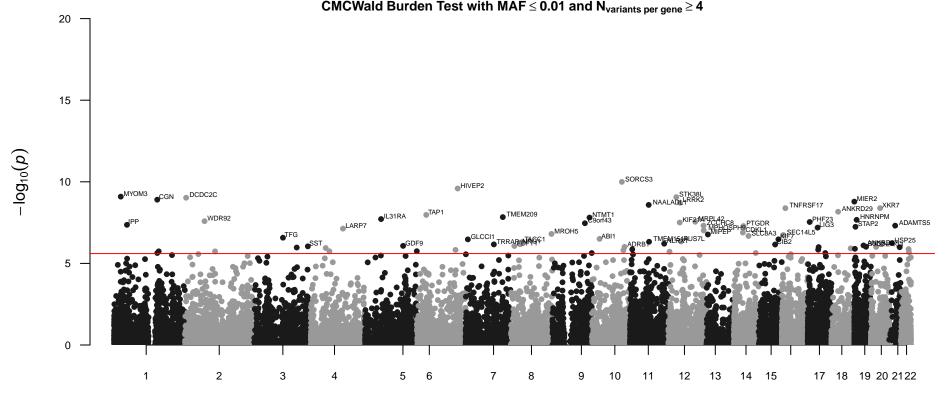
Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_ALL_PD_PHENOTYPES_CONTROL_2021

LOF (LOFTEE) and CADD > 20 (VEP)

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



Chromosome