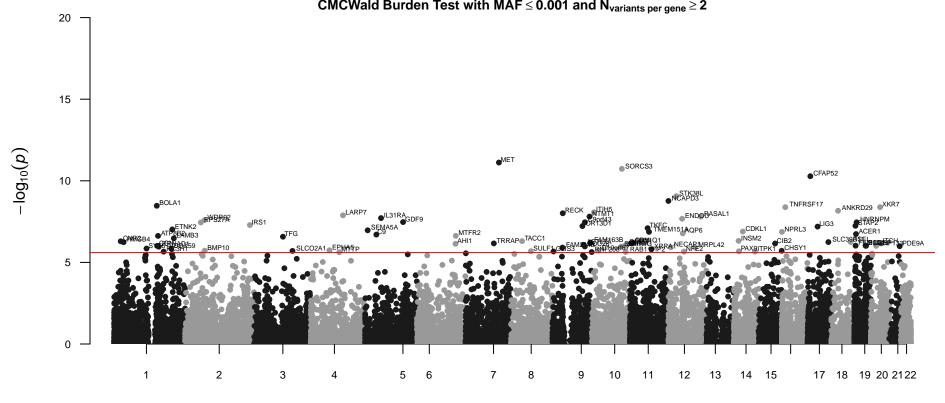
Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_ALL_PD_PHENOTYPES_CONTROL_2021

LOF (LOFTEE) and CADD > 20 (VEP)

CMCWald Burden Test with MAF \leq 0.001 and $N_{variants\ per\ gene} \geq$ 2



Chromosome