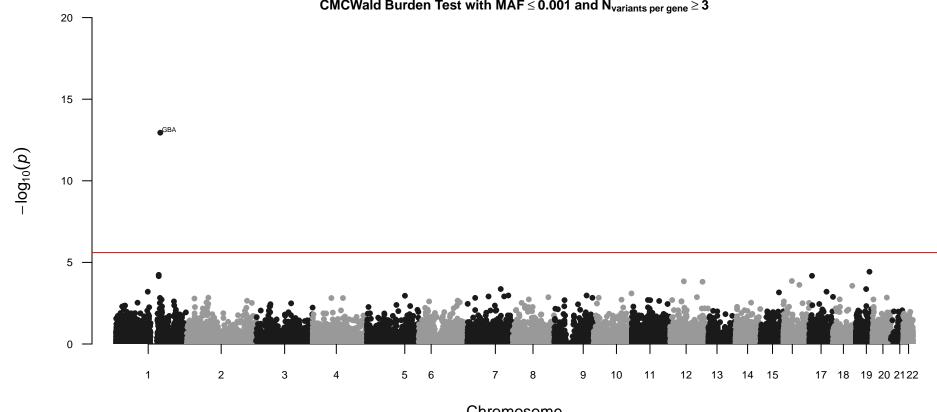
Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta_GNE_AMP_NIH_noRelateds_UKB_EXOM_PD_CASE_CONTROL_PARENT_PROXY_2021

Moderate or High Impact (SnpEff)

CMCWald Burden Test with MAF \leq 0.001 and $N_{variants\ per\ gene} \geq 3$



Chromosome