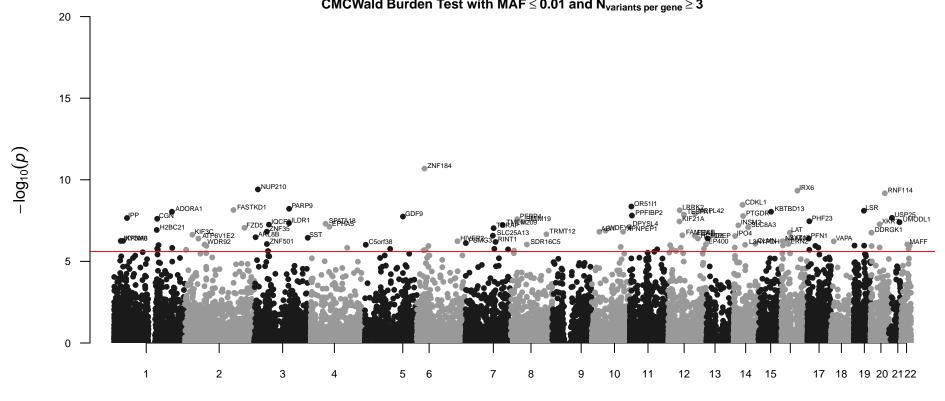
## Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta\_GNE\_AMP\_NIH\_noRelateds\_UKB\_EXOM\_ALL\_PD\_PHENOTYPES\_CONTROL\_2021

**Moderate or High Impact (SnpEff)** 

CMCWald Burden Test with MAF  $\leq$  0.01 and N<sub>variants per gene</sub>  $\geq$  3



Chromosome