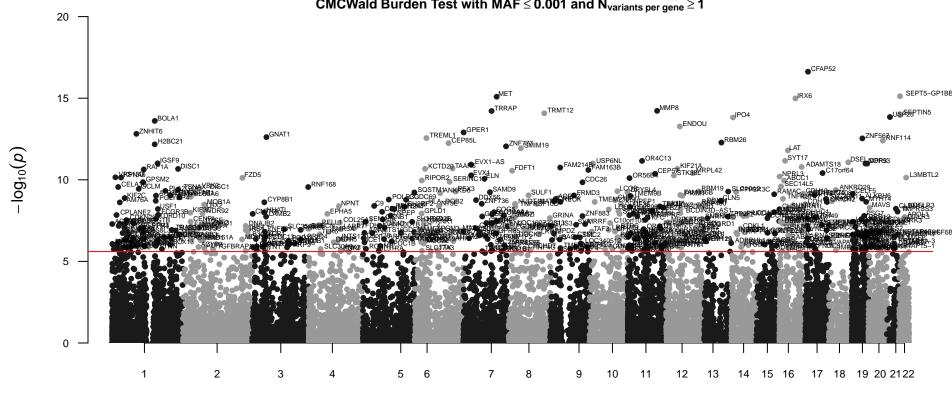
## Meta-analysis of AMP-PD x NIH WGS and UK Biobank WES

meta\_GNE\_AMP\_NIH\_noRelateds\_UKB\_EXOM\_PD\_CASE\_CONTROL\_PROXIES\_2021

Moderate or High Impact (SnpEff)

CMCWald Burden Test with MAF ≤ 0.001 and N<sub>variants per gene</sub> ≥ 1



Chromosome