

# 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)

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

