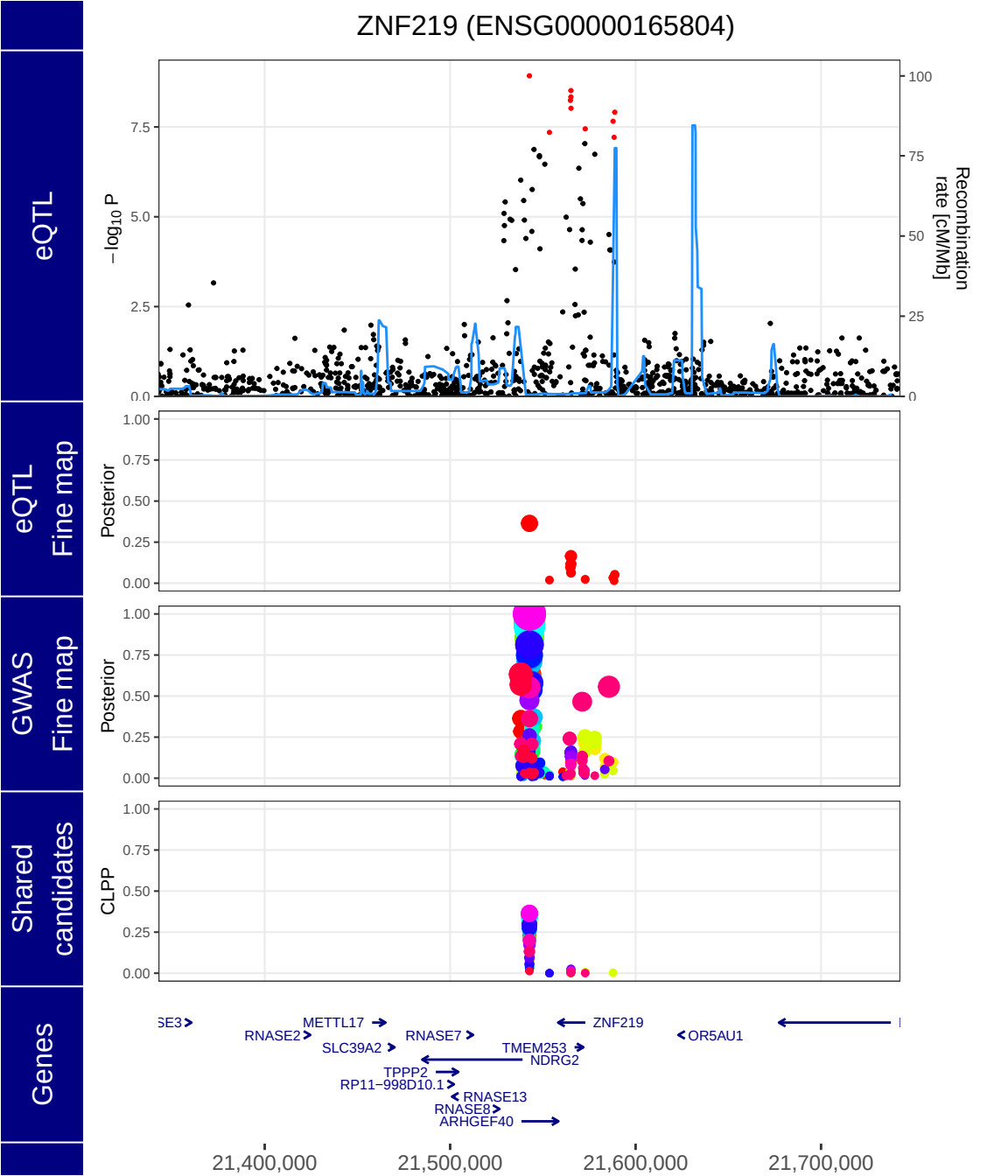
ZNF219 (ENSG00000165804)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
METTL17
RNASE2
SLC39A2
RNASE7
ZNF219
OR5AU1
TMEM253
NDRG2
TPPP2
RP11−998D10.1
RNASE13
RNASE8
ARHGEF40
21,400,000
21,500,000
21,600,000
21,700,000