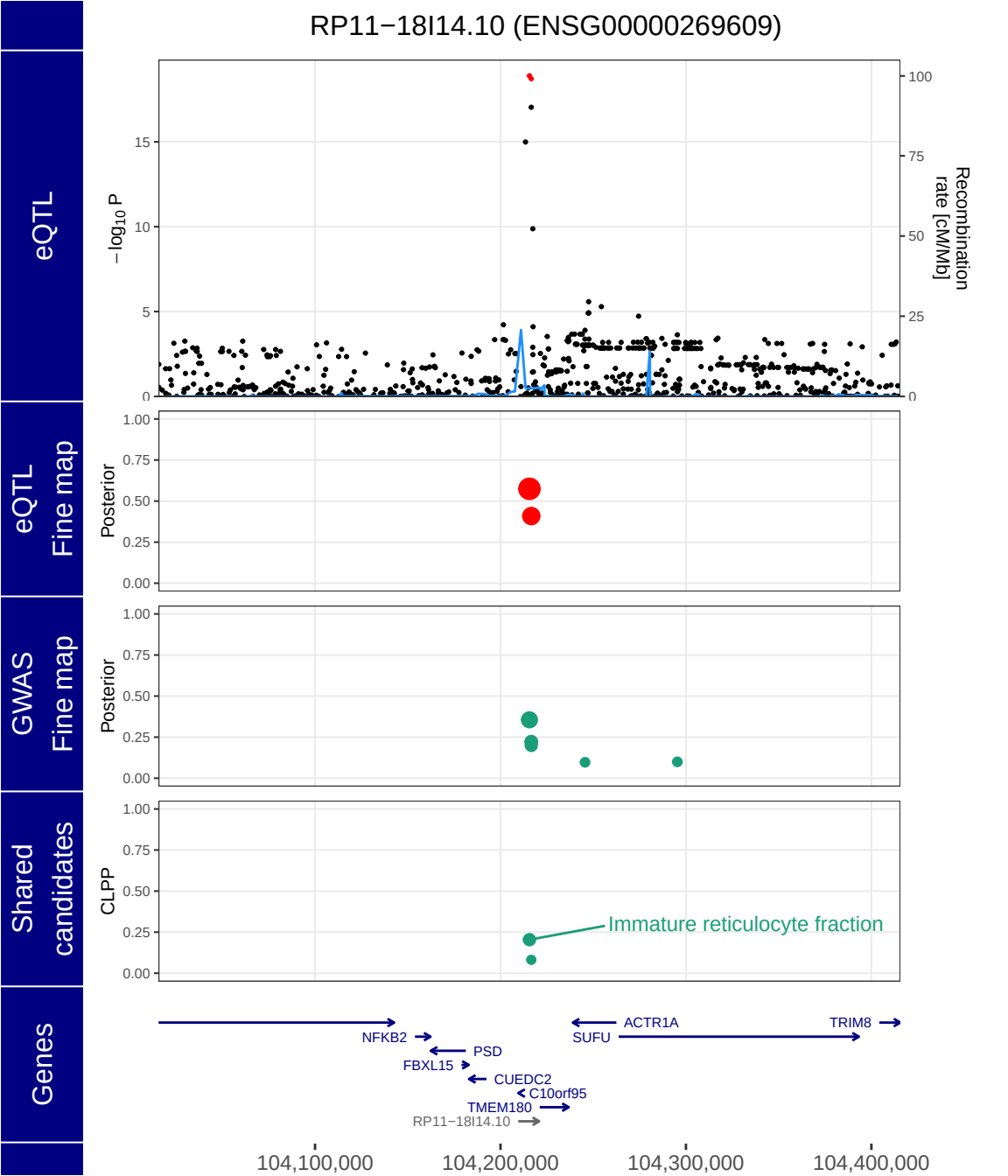
RP11−18I14.10 (ENSG00000269609)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Immature reticulocyte fraction
NFKB2
PSD
FBXL15
CUEDC2
C10orf95
TMEM180
RP11−18I14.10
ACTR1A
SUFU
TRIM8
104,100,000
104,200,000
104,300,000
104,400,000