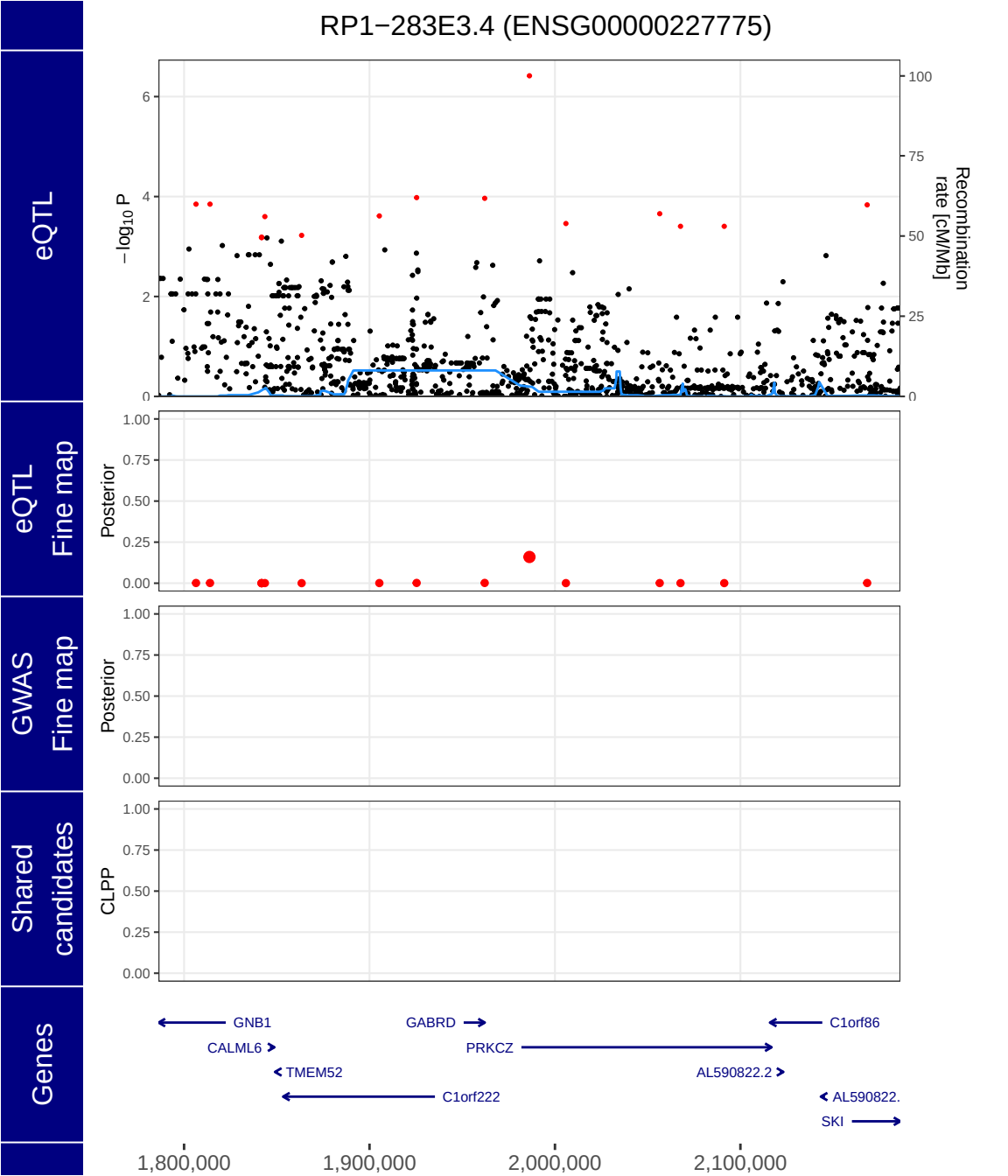
RP1−283E3.4 (ENSG00000227775)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
GNB1
CALML6
TMEM52
C1orf222
GABRD
PRKCZ
AL590822.2
C1orf86
AL590822.
SKI
1,800,000
1,900,000
2,000,000
2,100,000