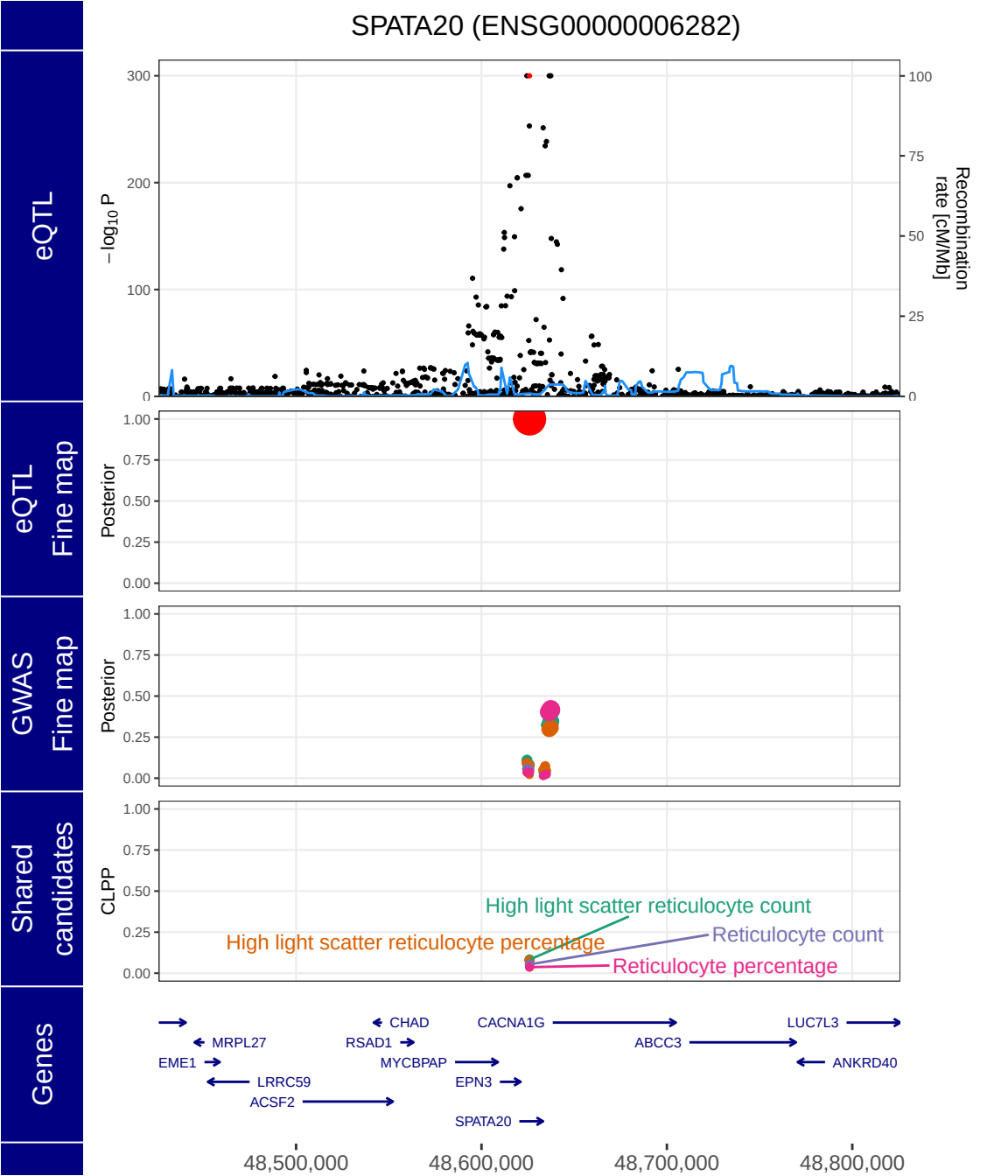
SPATA20 (ENSG00000006282)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
High light scatter reticulocyte count
Reticulocyte count
High light scatter reticulocyte percentage
Reticulocyte percentage
CHAD
CACNA1G
LUC7L3
MRPL27
RSAD1
ABCC3
EME1
MYCBPAP
ANKRD40
LRRC59
EPN3
ACSF2
SPATA20
48,500,000
48,600,000
48,700,000
48,800,000