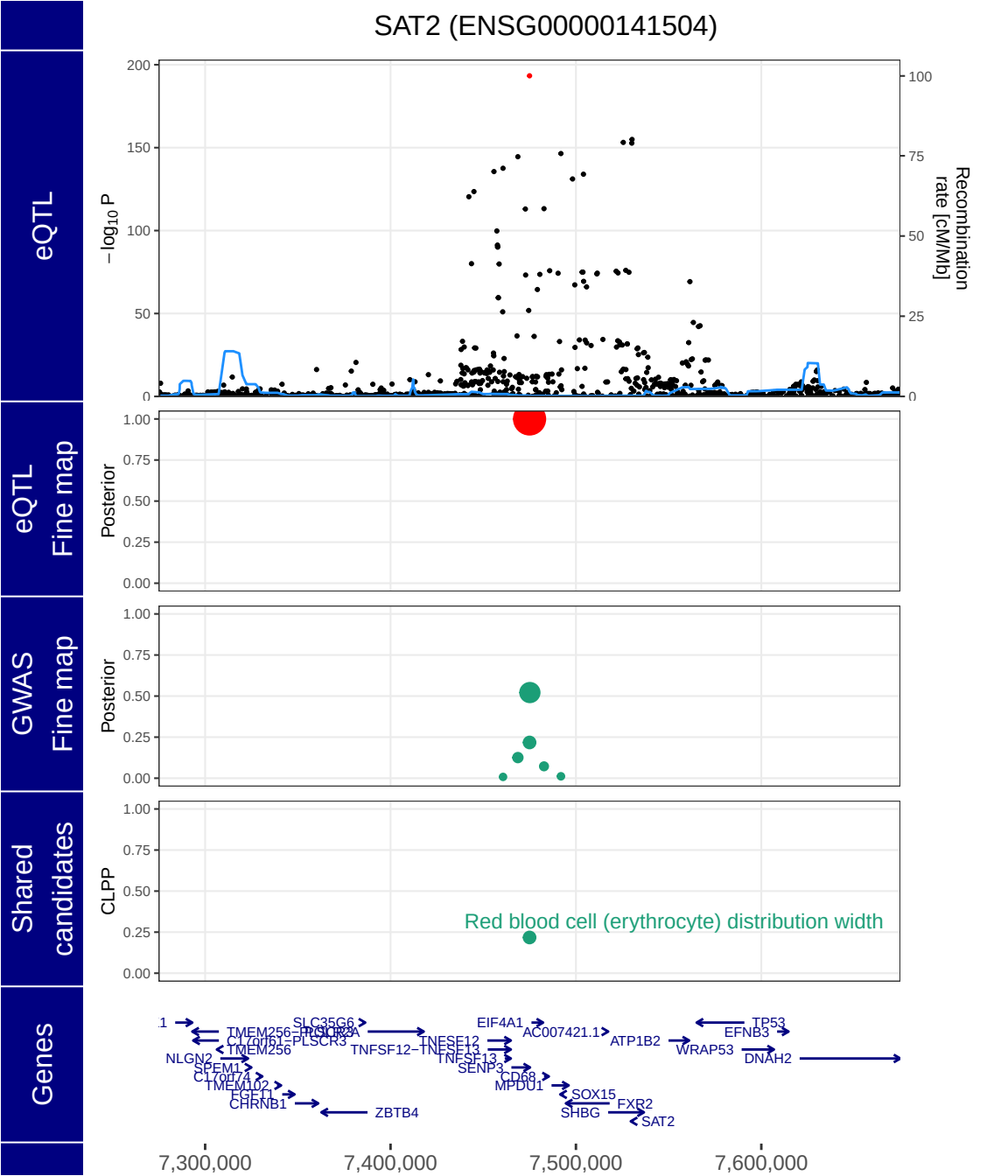
SAT2 (ENSG00000141504)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
Posterior
CLPP
Red blood cell (erythrocyte) distribution width
SLC35G6
TMEM256-PLSCR3
C17orf61
TMEM256
PLSCR3
NLGN2
SPEM1
C17orf74
TMEM102
EGFL7
CHRNB1
ZBTB4
TNFSF12-TNFSF13
TNFSF12
TNFSF13
TNFSF13
SENP3
EIF4A1
AC007421.1
ATP1B2
CD68
MPDU1
SOX15
FXR2
SHBG
SAT2
TP53
EFNB3
WRAP53
DNAH2
7,300,000
7,400,000
7,500,000
7,600,000