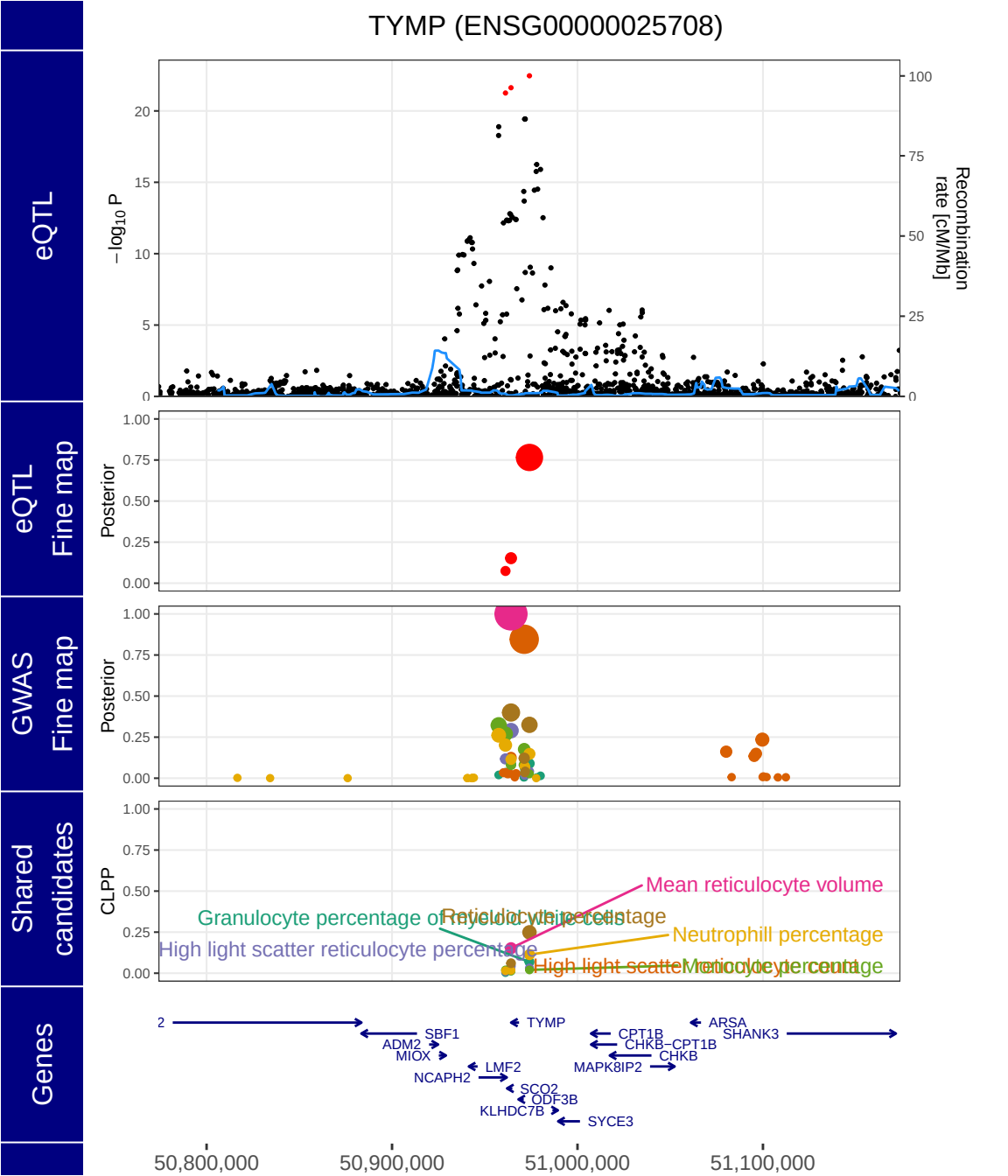
TYMP (ENSG00000025708)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Mean reticulocyte volume
Granulocyte percentage of myeloid white cells
Reticulocyte percentage
Neutrophill percentage
High light scatter reticulocyte percentage
High light scatter reticulocyte count
Monocyte percentage
TYMP
ARSA
SHANK3
SBF1
CPT1B
CHKB−CPT1B
ADM2
MIOX
CHKB
MAPK8IP2
LMF2
NCAPH2
SCO2
ODF3B
KLHDC7B
SYCE3
50,800,000
50,900,000
51,000,000
51,100,000