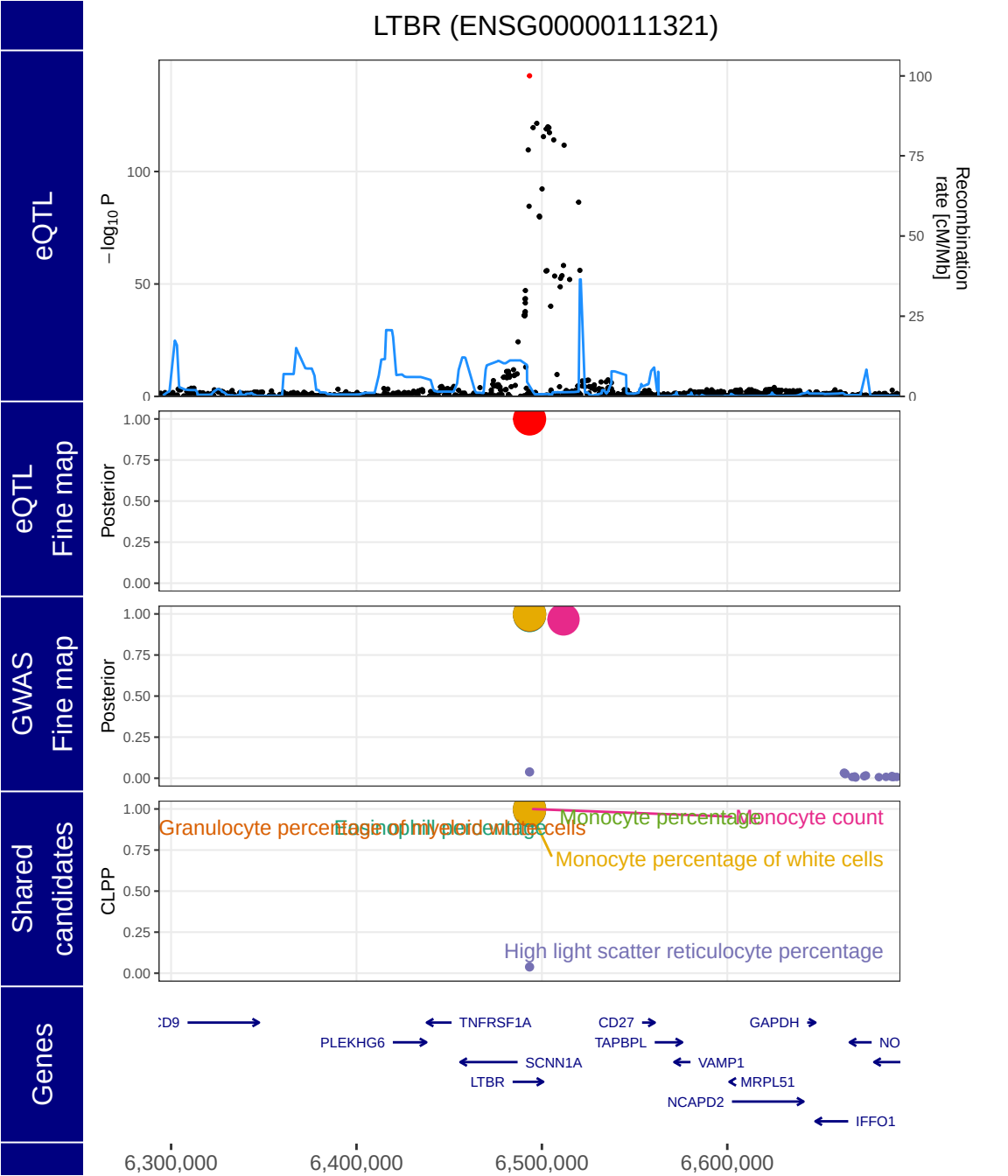
LTBR (ENSG00000111321)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Granulocyte percentage of myeloid white cells
Eosinophil percentage
Monocyte percentage
Monocyte count
Monocyte percentage of white cells
High light scatter reticulocyte percentage
TNFRSF1A
CD27
GAPDH
PLEKHG6
TAPBPL
SCNN1A
VAMP1
LTBR
MRPL51
NCAPD2
IFFO1
6,300,000
6,400,000
6,500,000
6,600,000