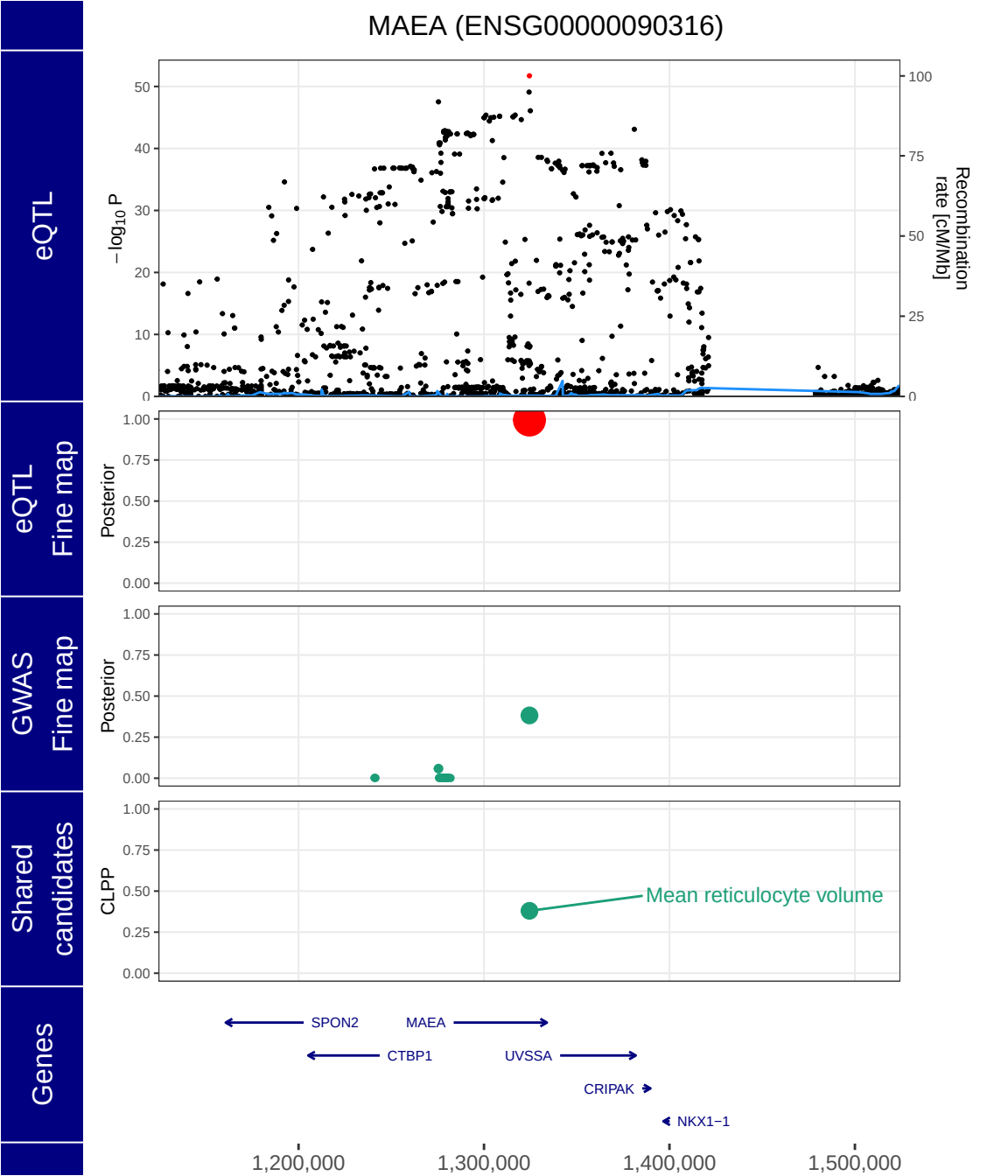
MAEA (ENSG00000090316)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Mean reticulocyte volume
SPON2
MAEA
CTBP1
UVSSA
CRIPAK
NKX1−1
1,200,000
1,300,000
1,400,000
1,500,000