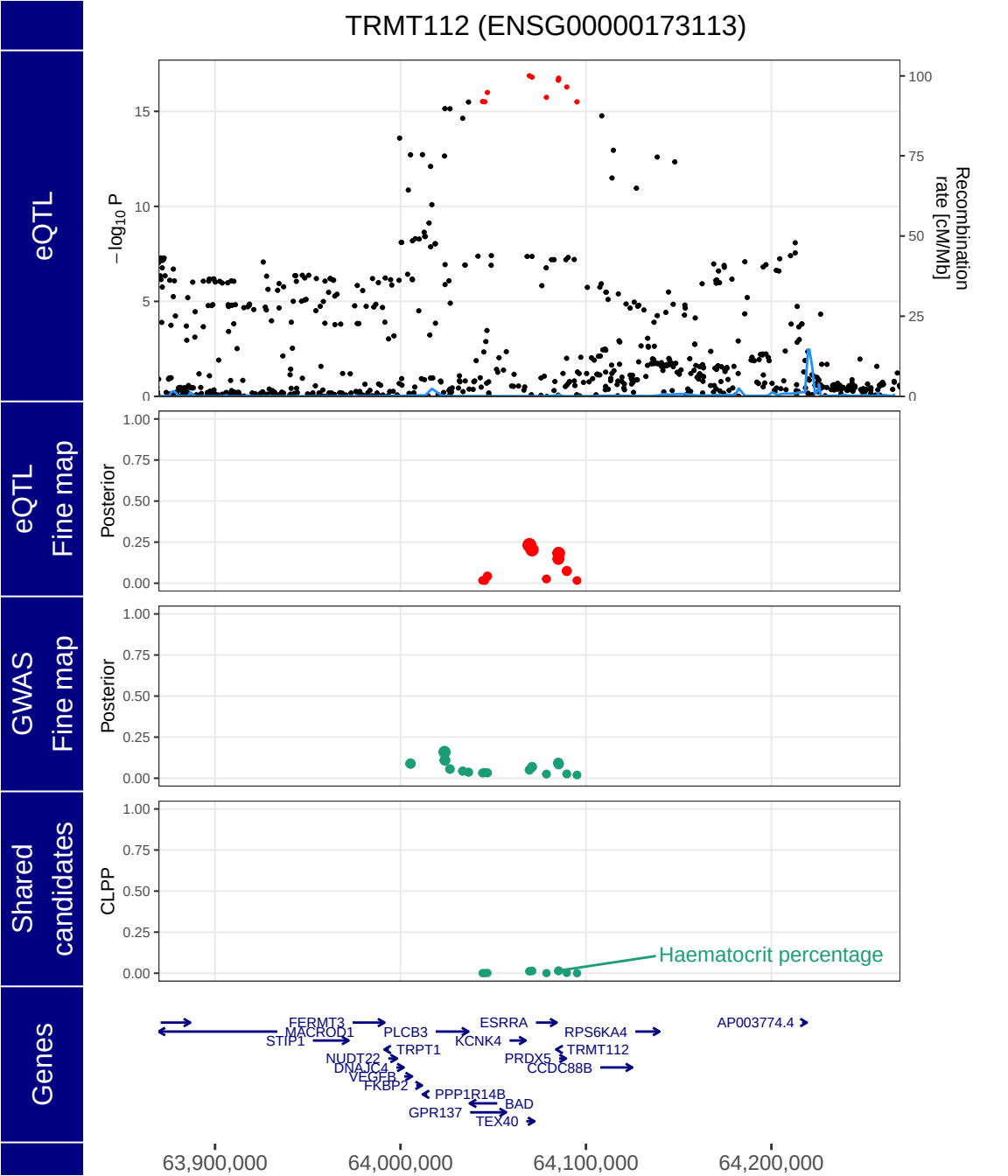
TRMT112 (ENSG00000173113)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Haematocrit percentage
FERMT3
MACROD1
STIP1
PLCB3
TRPT1
NUDT22
DNAJC4
VEGFB
FKBP2
PPP1R14B
GPR137
BAD
TEX40
ESRRA
KCNK4
RPS6KA4
TRMT112
PRDX5
CCDC88B
AP003774.4
63,900,000
64,000,000
64,100,000
64,200,000