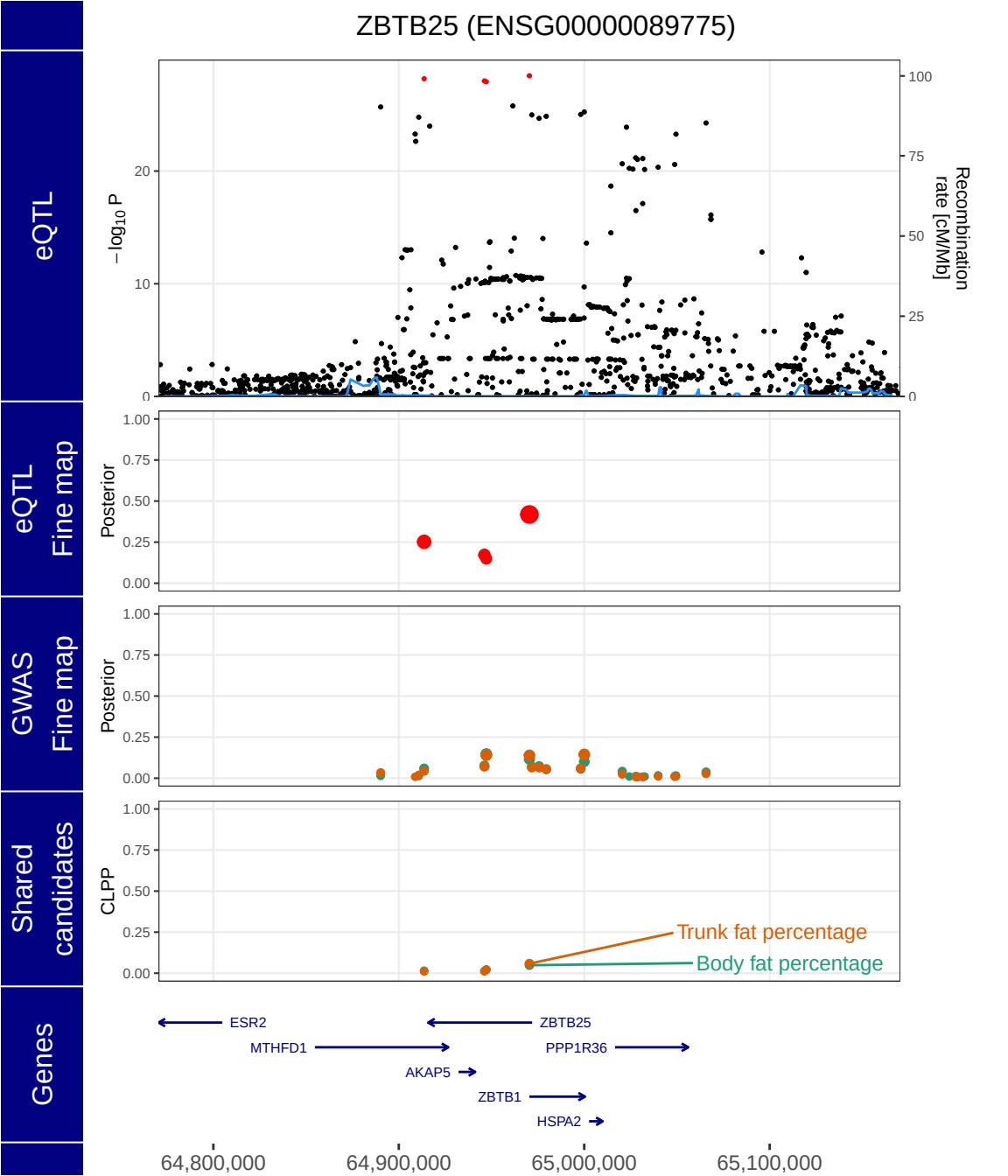
ZBTB25 (ENSG00000089775)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Trunk fat percentage
Body fat percentage
ESR2
ZBTB25
MTHFD1
PPP1R36
AKAP5
ZBTB1
HSPA2
64,800,000
64,900,000
65,000,000
65,100,000