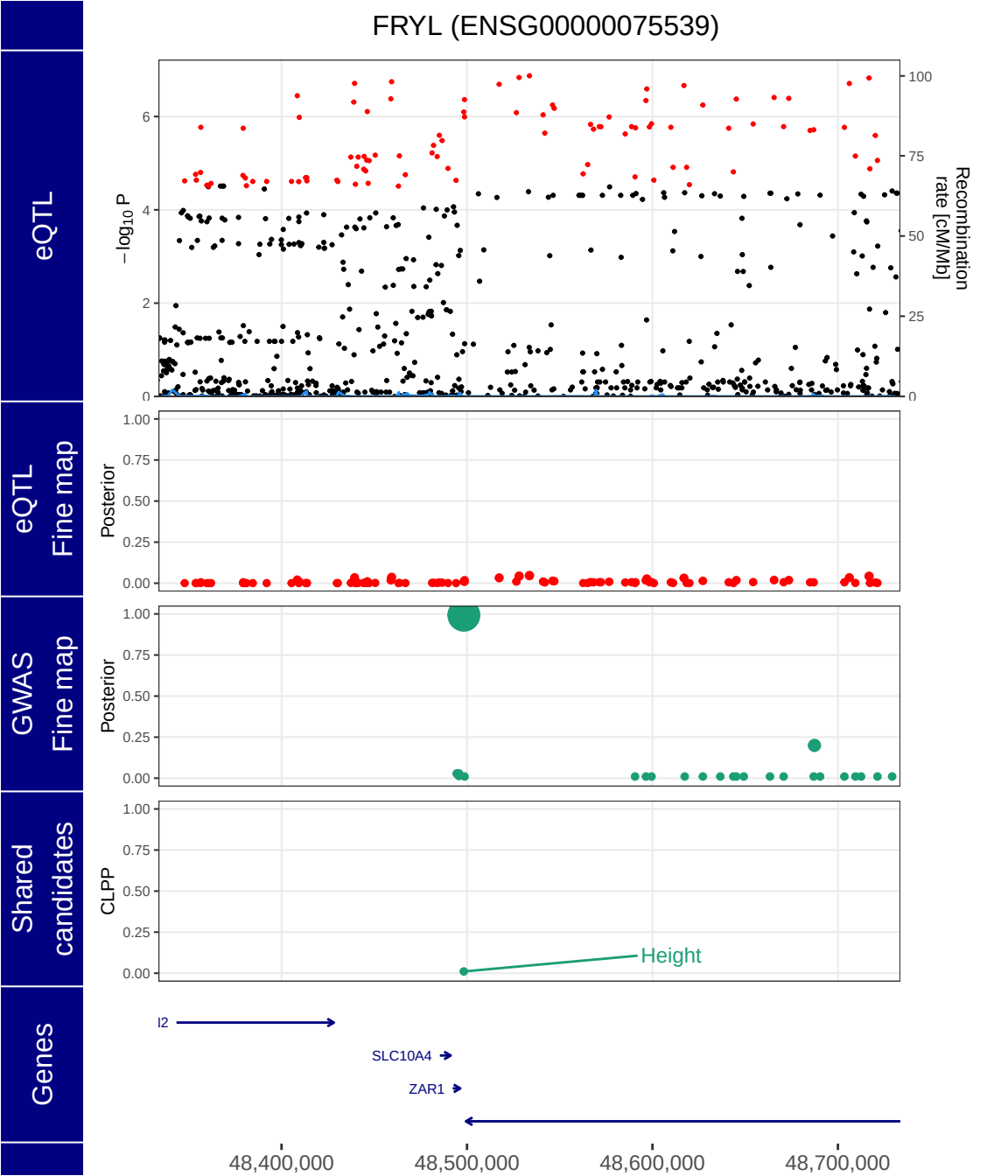
FRYL (ENSG00000075539)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Height
SLC10A4
ZAR1
48,400,000
48,500,000
48,600,000
48,700,000