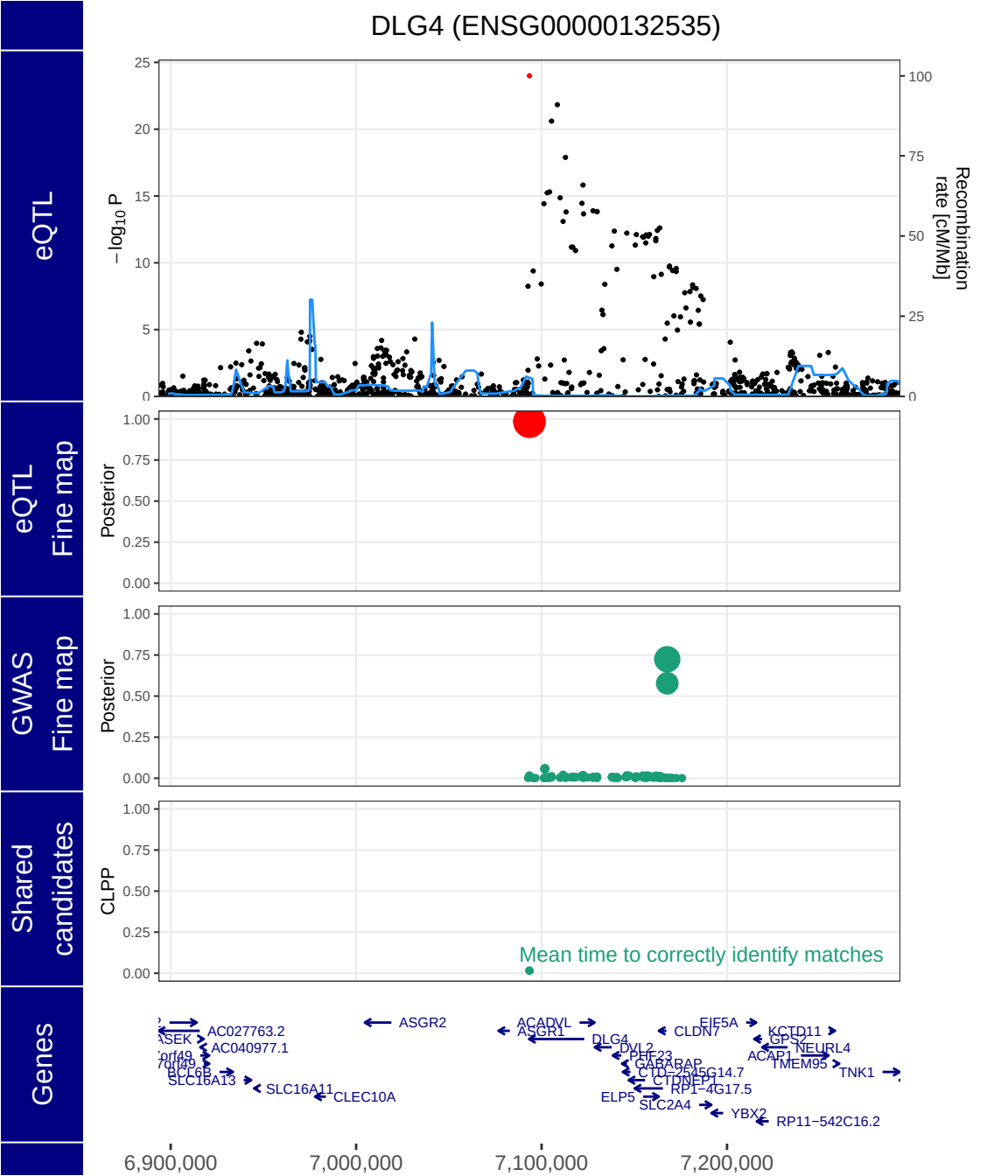
DLG4 (ENSG00000132535)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Mean time to correctly identify matches
AC027763.2
AC040977.1
SLC16A13
SLC16A11
CLEC10A
ASGR2
ACADVL
ASGR1
DLG4
DVL2
PHF23
GARARAP
C1D−2545G14.7
CTDNEP1
RP1−4G17.5
ELP5
SLC2A4
YBX2
RP11−542C16.2
EIF5A
CLDN7
KCTD11
GPS2
NEURL4
ACAP1
TMEM95
TNK1
6,900,000
7,000,000
7,100,000
7,200,000