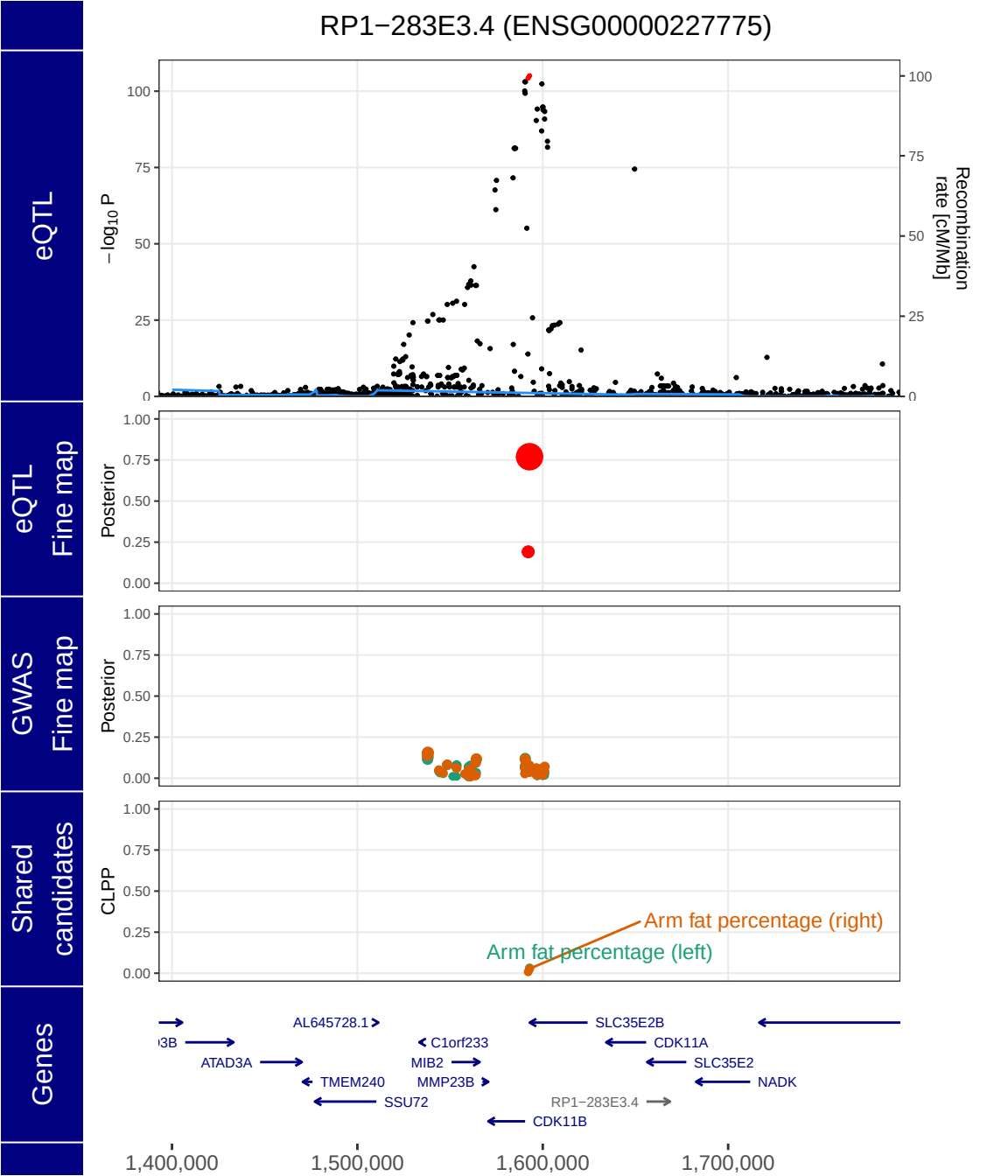
RP1−283E3.4 (ENSG00000227775)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Arm fat percentage (right)
Arm fat percentage (left)
AL645728.1
SLC35E2B
C1orf233
CDK11A
ATAD3A
MIB2
SLC35E2
TMEM240
MMP23B
NADK
SSU72
RP1−283E3.4
CDK11B
1,400,000
1,500,000
1,600,000
1,700,000