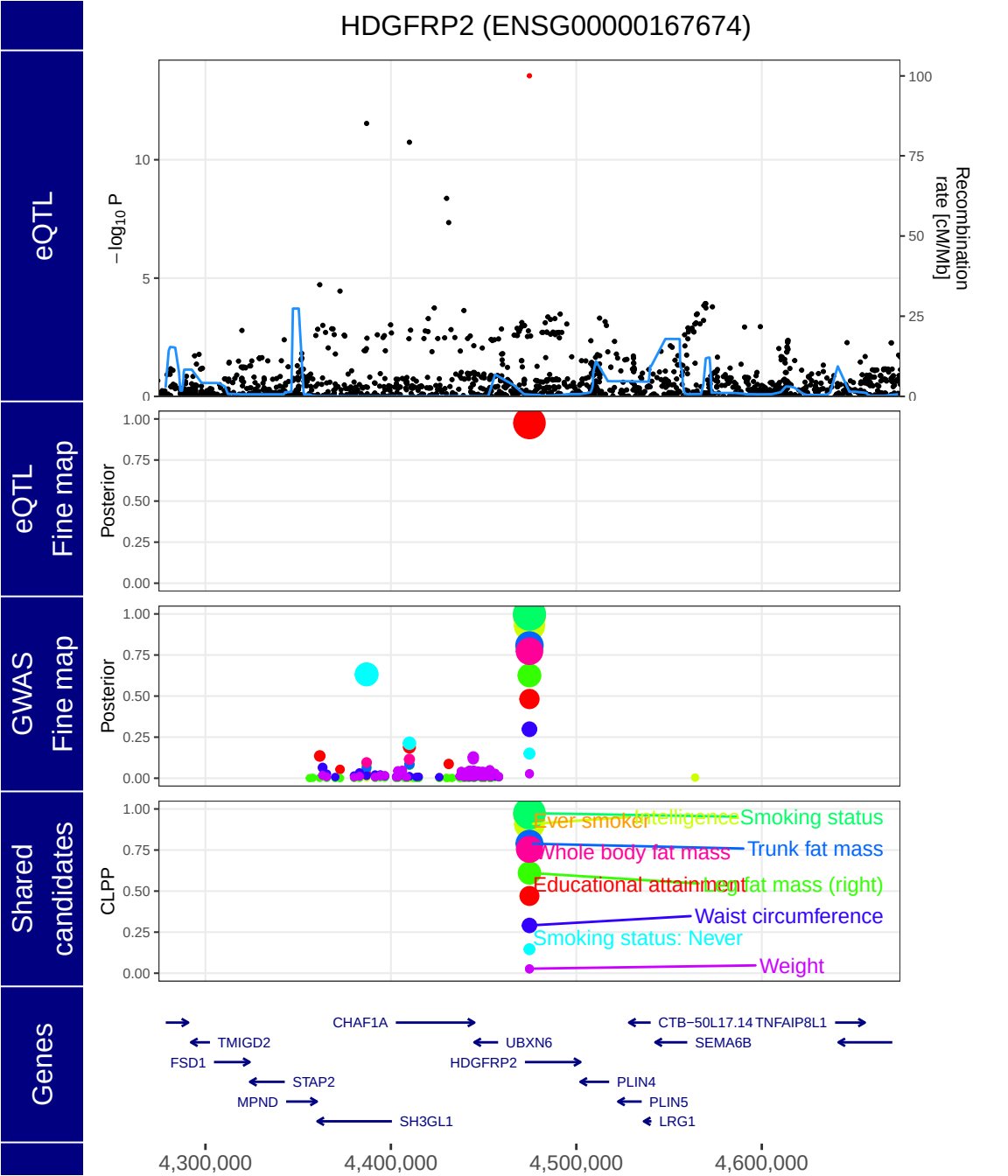
HDGFRP2 (ENSG00000167674)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Ever smoker
Intelligence
Smoking status
Whole body fat mass
Trunk fat mass
Educational attainment
Leg fat mass (right)
Waist circumference
Smoking status: Never
Weight
CHAF1A
TMIGD2
FSD1
STAP2
MPND
SH3GL1
UBXN6
HDGFRP2
CTB−50L17.14
TNFAIP8L1
SEMA6B
PLIN4
PLIN5
LRG1
4,300,000
4,400,000
4,500,000
4,600,000