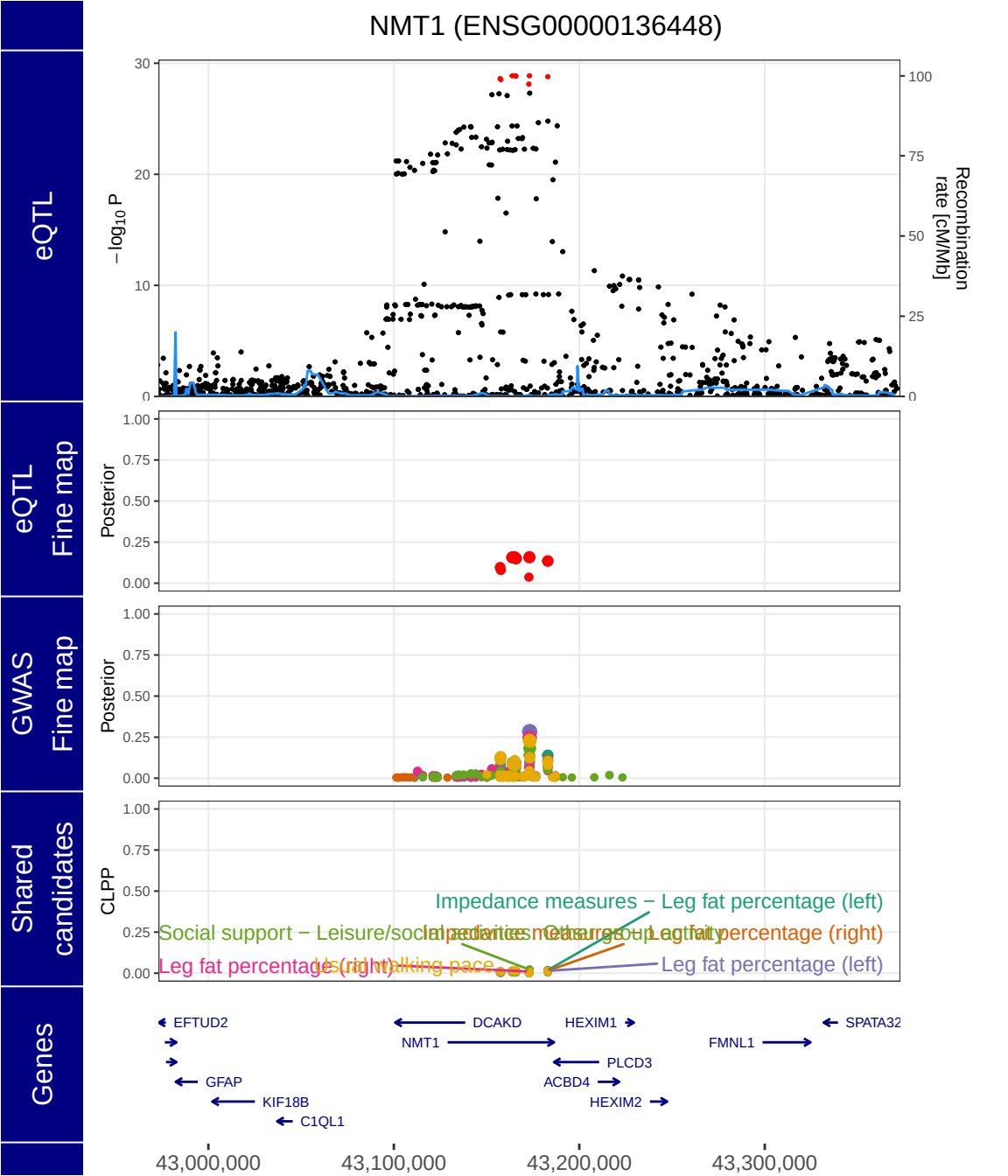
NMT1 (ENSG00000136448)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Impedance measures − Leg fat percentage (left)
Social support − Leisure/social activities
Leg fat percentage (right)
Leg fat percentage (left)
EFTUD2
DCAKD
HEXIM1
SPATA32
NMT1
FMNL1
PLCD3
GFAP
ACBD4
KIF18B
HEXIM2
C1QL1
43,000,000
43,100,000
43,200,000
43,300,000