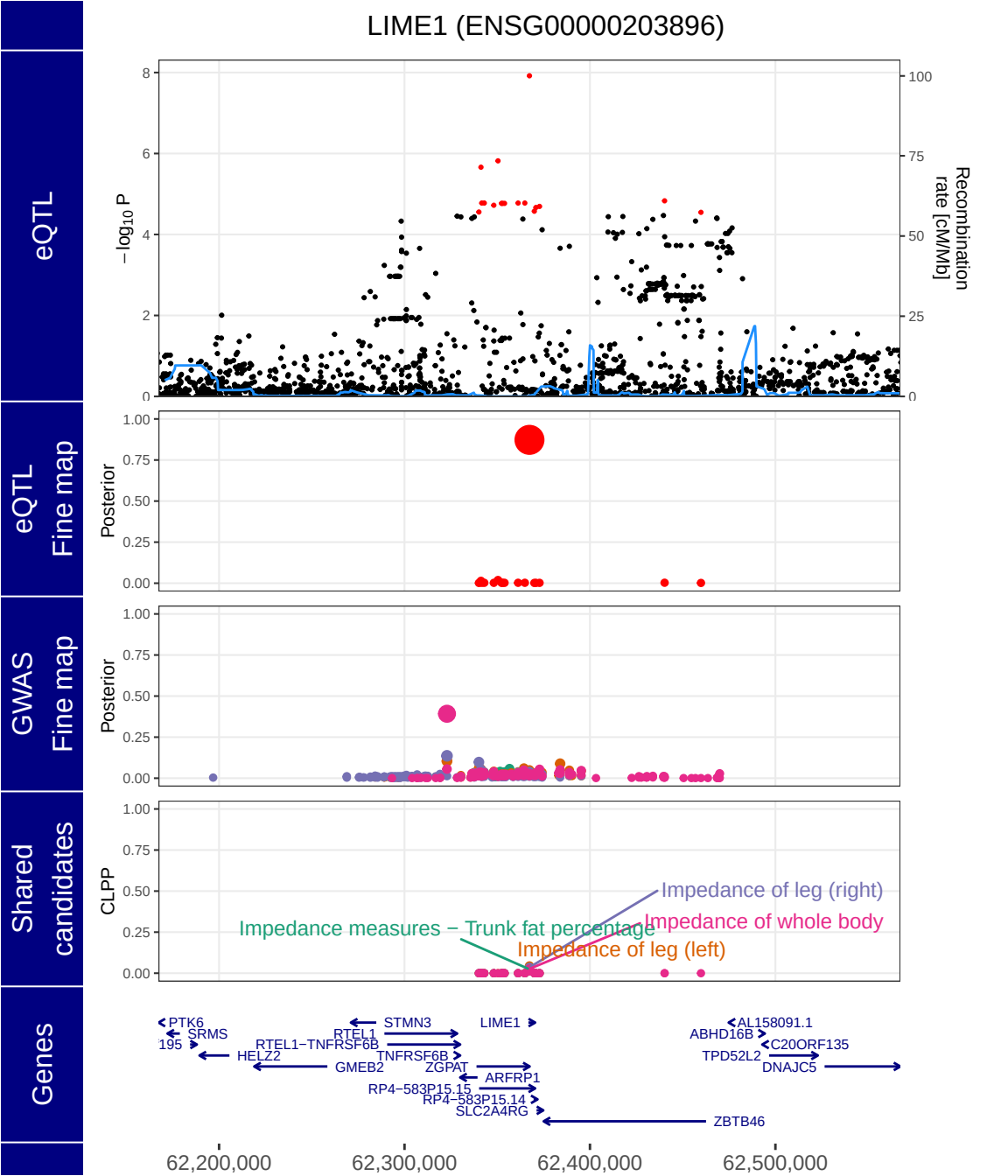
LIME1 (ENSG00000203896)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Impedance of leg (right)
Impedance of whole body
Impedance measures − Trunk fat percentage
Impedance of leg (left)
PTK6
SRMS
195
HELZ2
RTEL1−TNFRSF6B
GMEB2
STMN3
RTEL1
TNFRSF6B
ZGPAT
LIME1
ARFRP1
RP4−583P15.15
RP4−583P15.14
SLC2A4RG
ZBTB46
AL158091.1
ABHD16B
C20ORF135
TPD52L2
DNAJC5
62,200,000
62,300,000
62,400,000
62,500,000