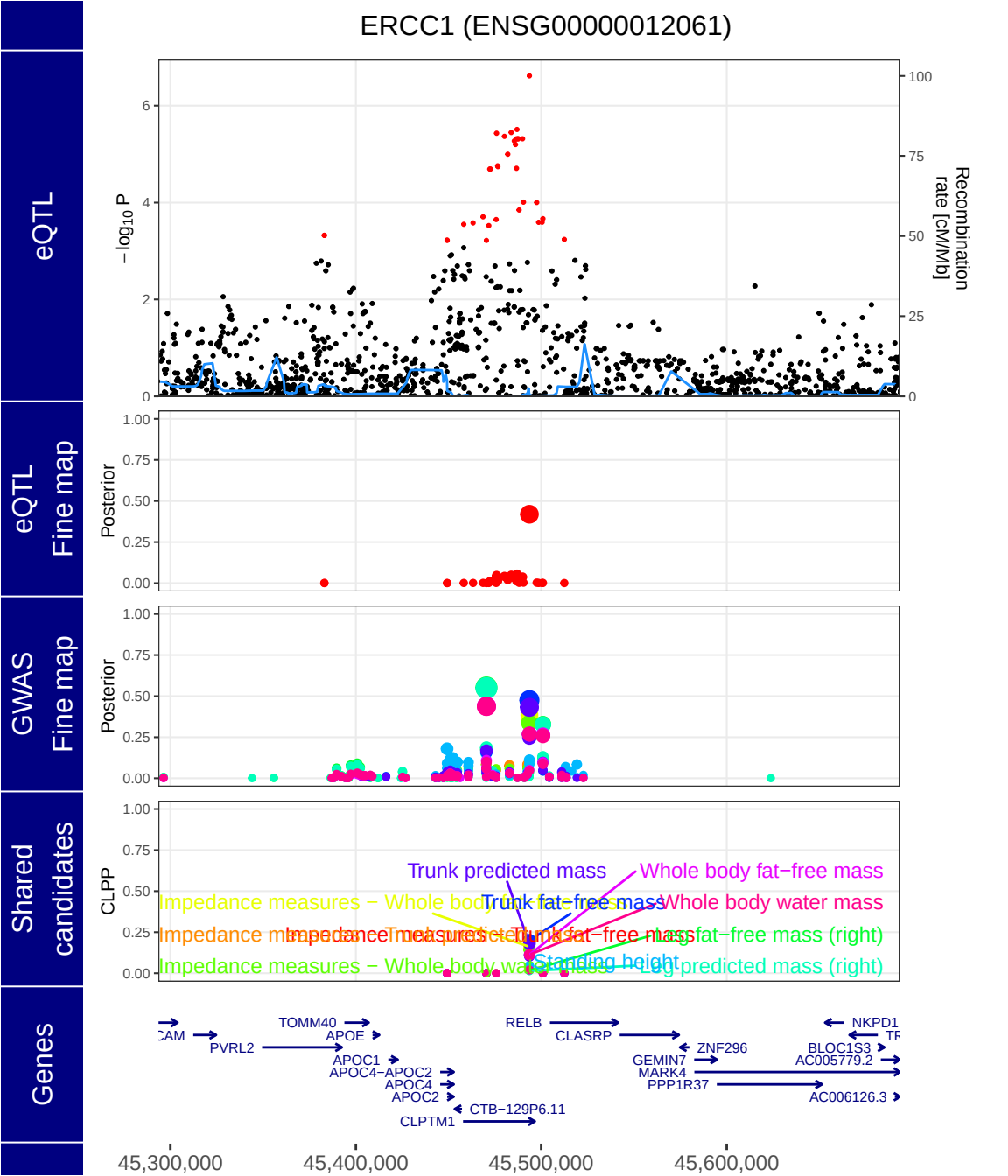
ERCC1 (ENSG00000012061)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Trunk predicted mass
Whole body fat−free mass
Trunk fat−free mass
Whole body water mass
Standing height
Leg predicted mass (right)
TOMM40
APOE
PVRL2
APOC1
APOC4−APOC2
APOC4
APOC2
CTB−129P6.11
CLPTM1
RELB
CLASRP
ZNF296
GEMIN7
MARK4
PPP1R37
NKPD1
BLOC1S3
AC005779.2
AC006126.3
45,300,000
45,400,000
45,500,000
45,600,000