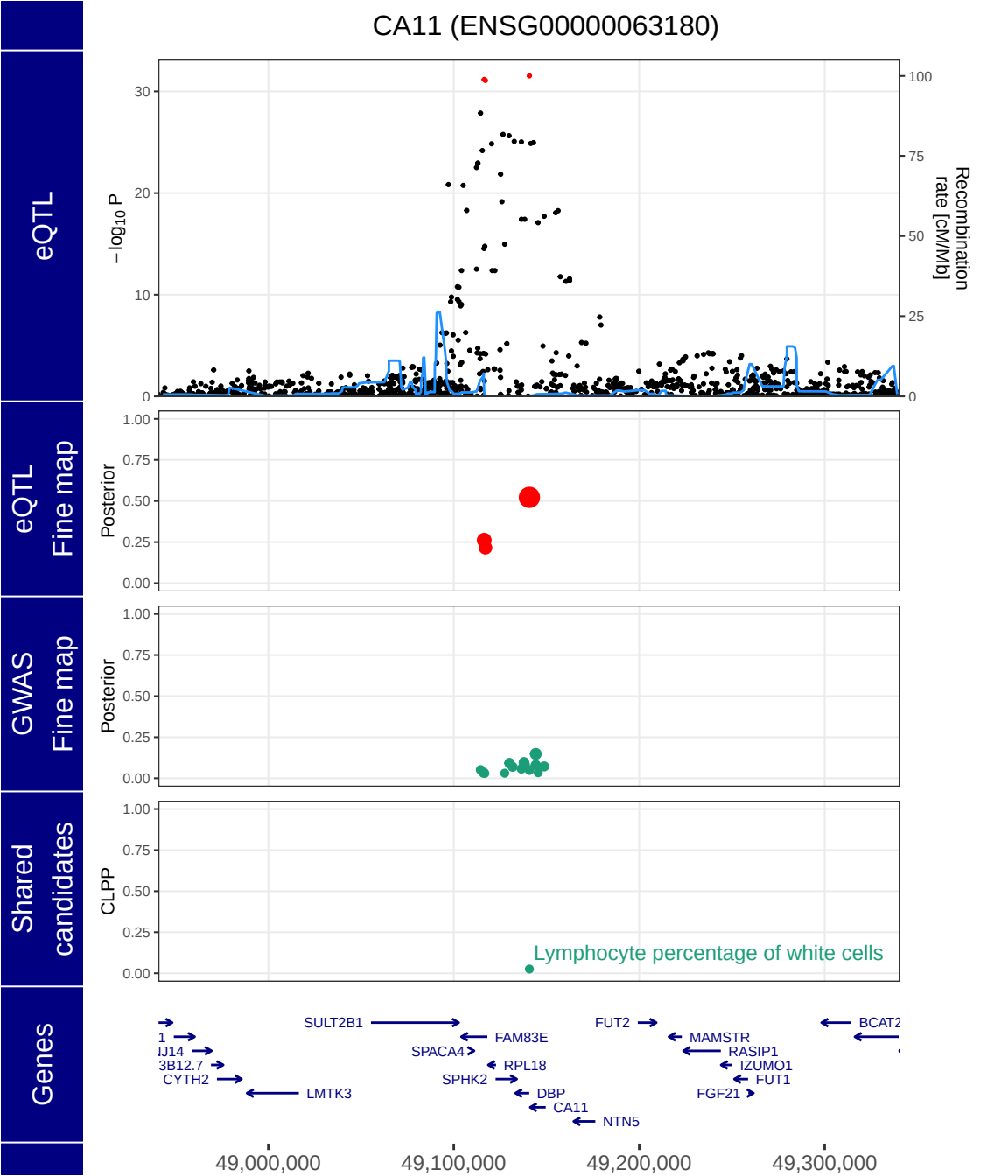
CA11 (ENSG00000063180)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Lymphocyte percentage of white cells
SULT2B1
FAM83E
SPACA4
RPL18
SPHK2
DBP
CA11
NTN5
LMTK3
CYTH2
3B12.7
IJ14
1
FUT2
MAMSTR
RASIP1
IZUMO1
FUT1
FGF21
BCAT2
49,000,000
49,100,000
49,200,000
49,300,000