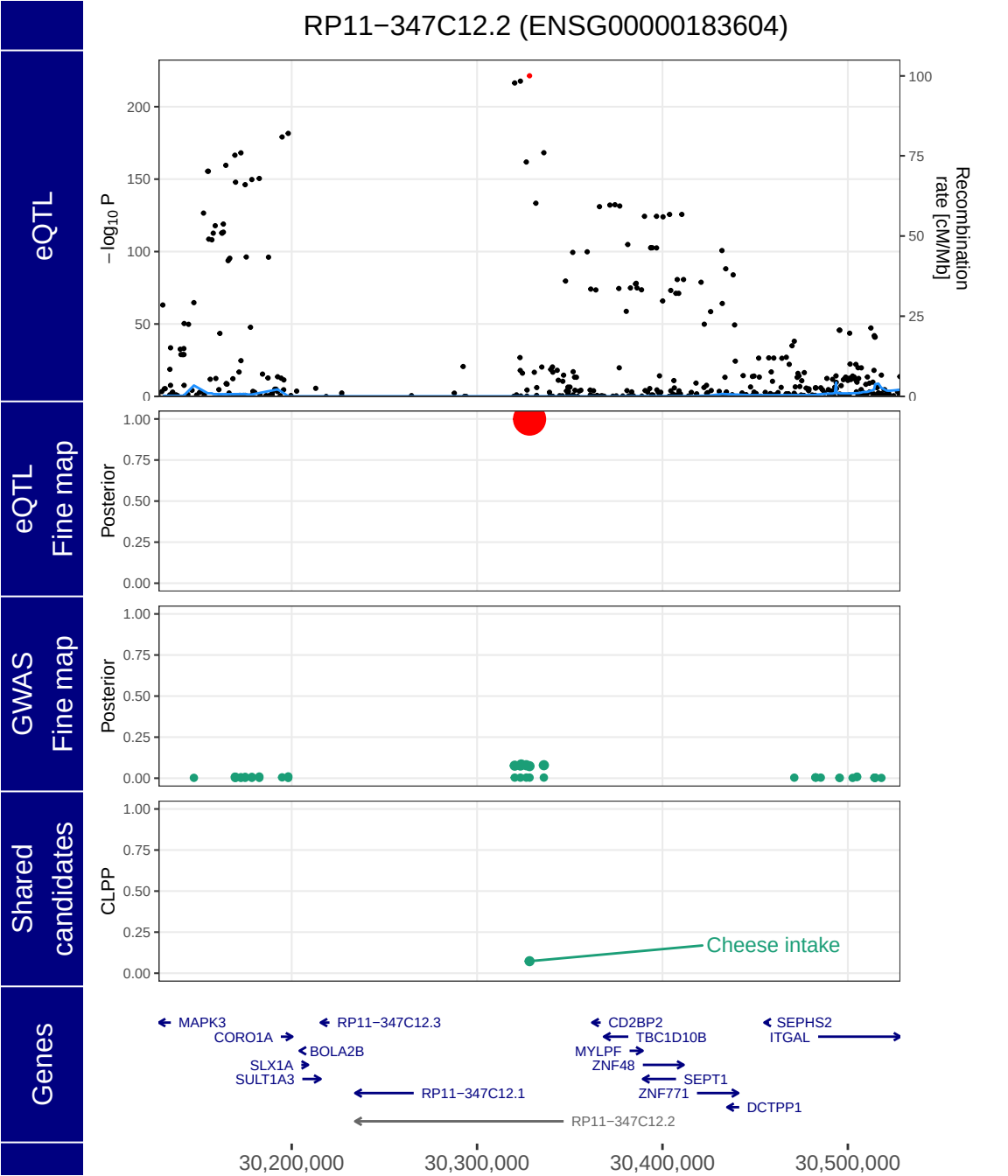
RP11−347C12.2 (ENSG00000183604)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
CLPP
Cheese intake
MAPK3
CORO1A
BOLA2B
SLX1A
SULT1A3
RP11−347C12.3
RP11−347C12.1
RP11−347C12.2
CD2BP2
TBC1D10B
MYLPF
ZNF48
SEPT1
ZNF771
DCTPP1
SEPHS2
ITGAL
30,200,000
30,300,000
30,400,000
30,500,000