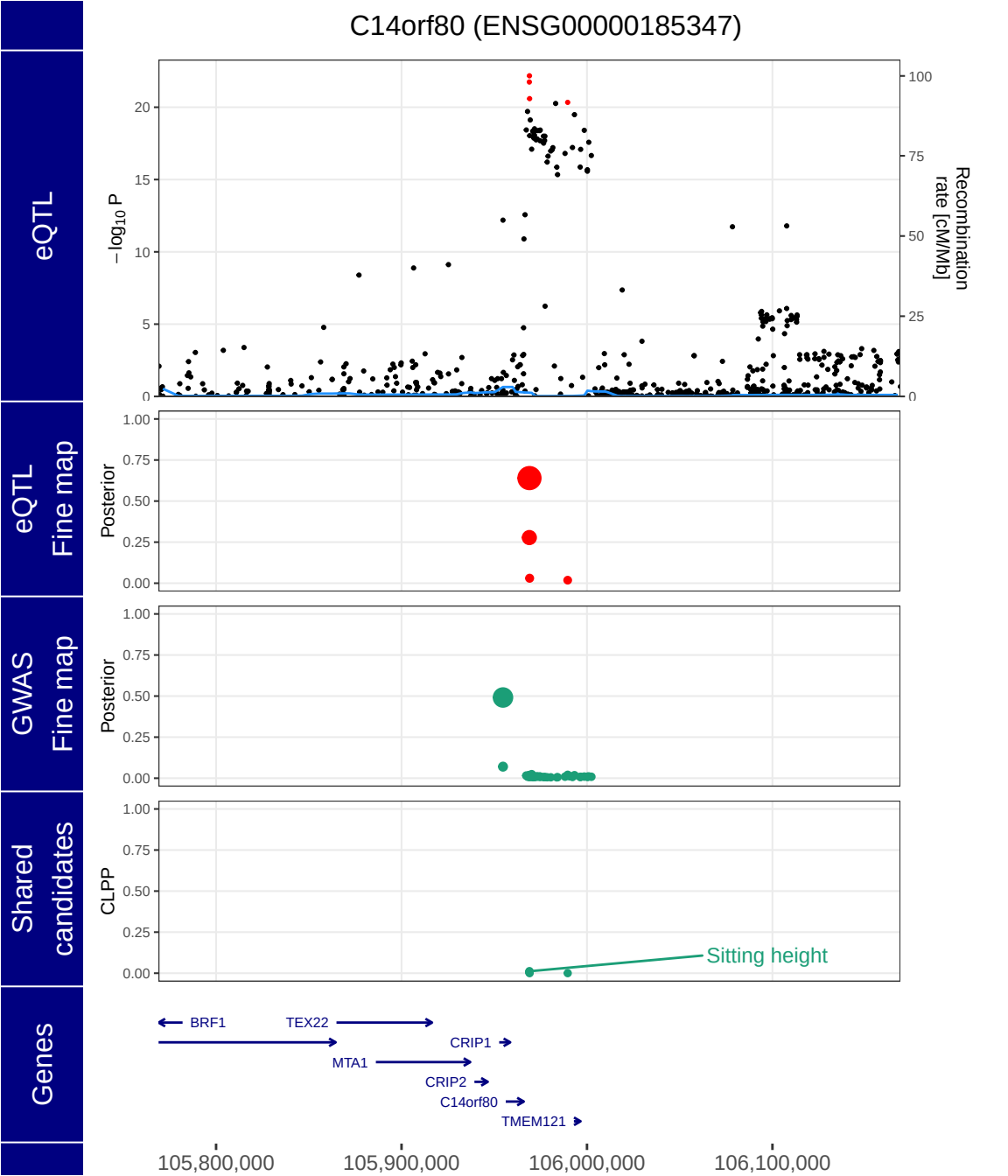
C14orf80 (ENSG00000185347)
eQTL
eQTL Fine map
GWAS Fine map
Shared candidates
Genes
−log10 P
Recombination rate [cM/Mb]
Posterior
Posterior
CLPP
Sitting height
BRF1
TEX22
CRIP1
MTA1
CRIP2
C14orf80
TMEM121
105,800,000
105,900,000
106,000,000
106,100,000