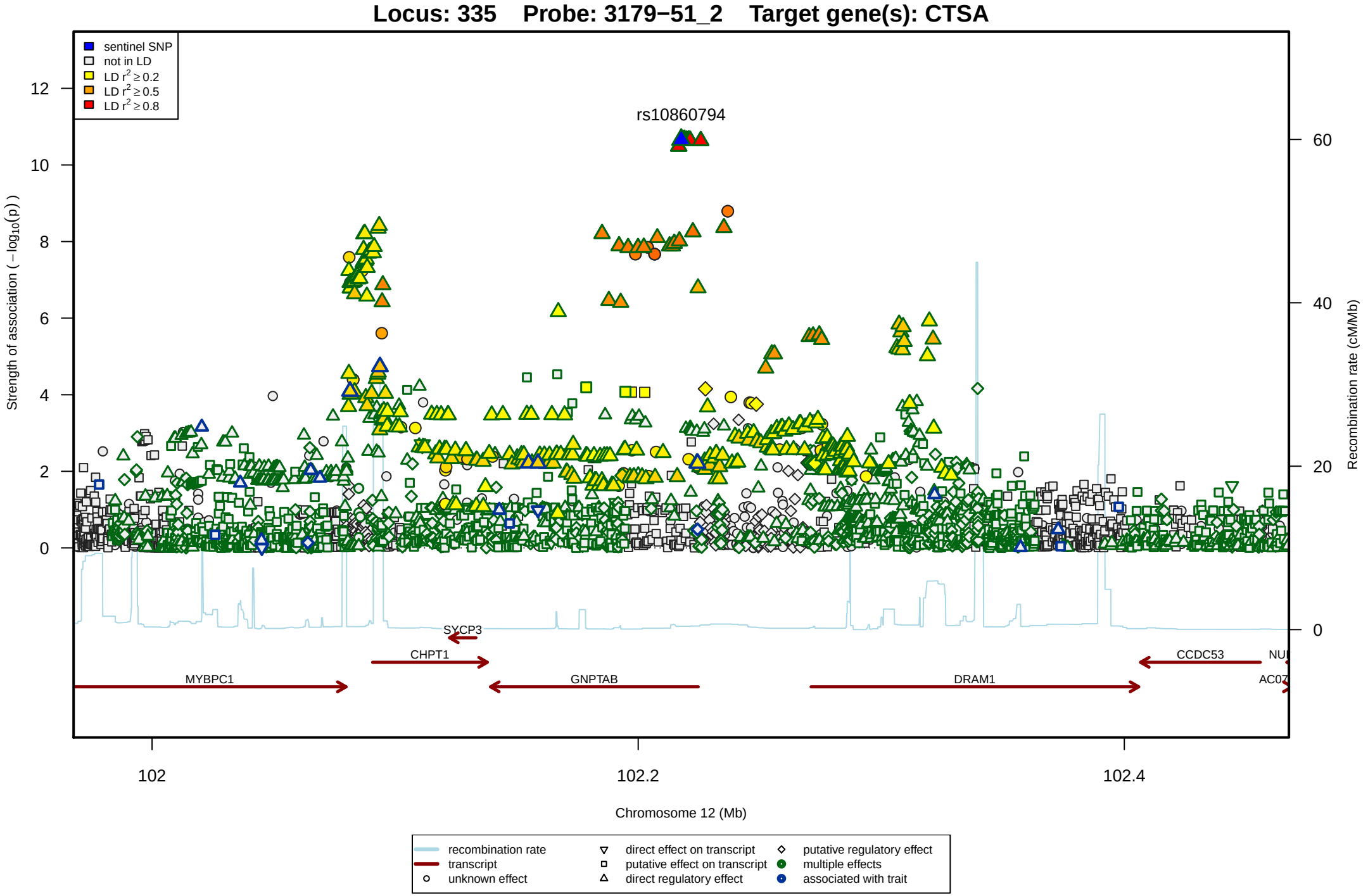

Locus: 335 Probe: 3179−51_2 Target gene(s): CTSA
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs10860794
Strength of association (−log10(p))
Recombination rate (cM/Mb)
SYCP3
CHPT1
MYBPC1
GNPTAB
DRAM1
CCDC53
Chromosome 12 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait