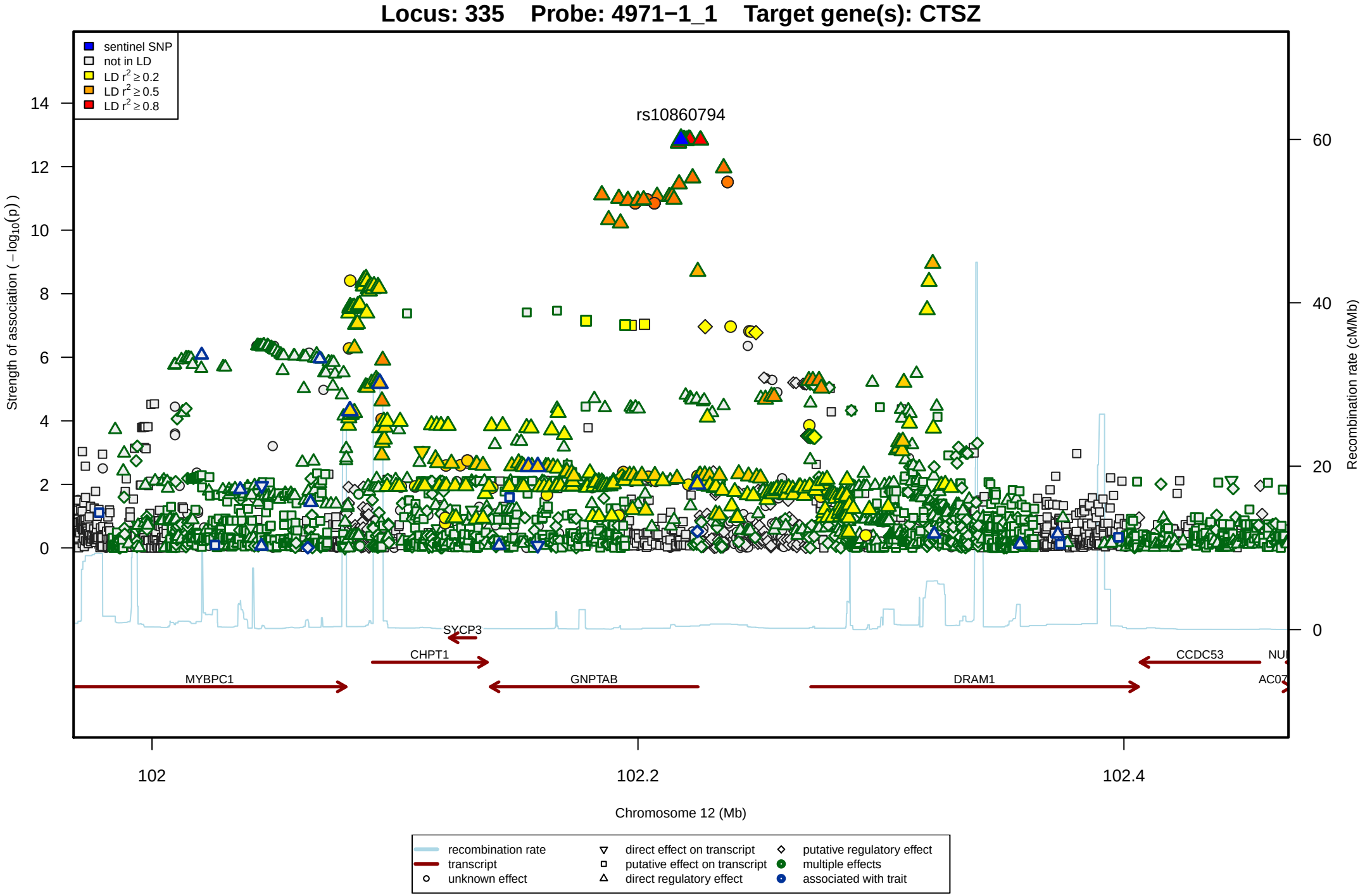

Locus: 335 Probe: 4971−1_1 Target gene(s): CTSZ
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs10860794
Strength of association (−log10(p))
Recombination rate (cM/Mb)
SYCP3
CHPT1
MYBPC1
GNPTAB
DRAM1
CCDC53
102
102.2
102.4
Chromosome 12 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait