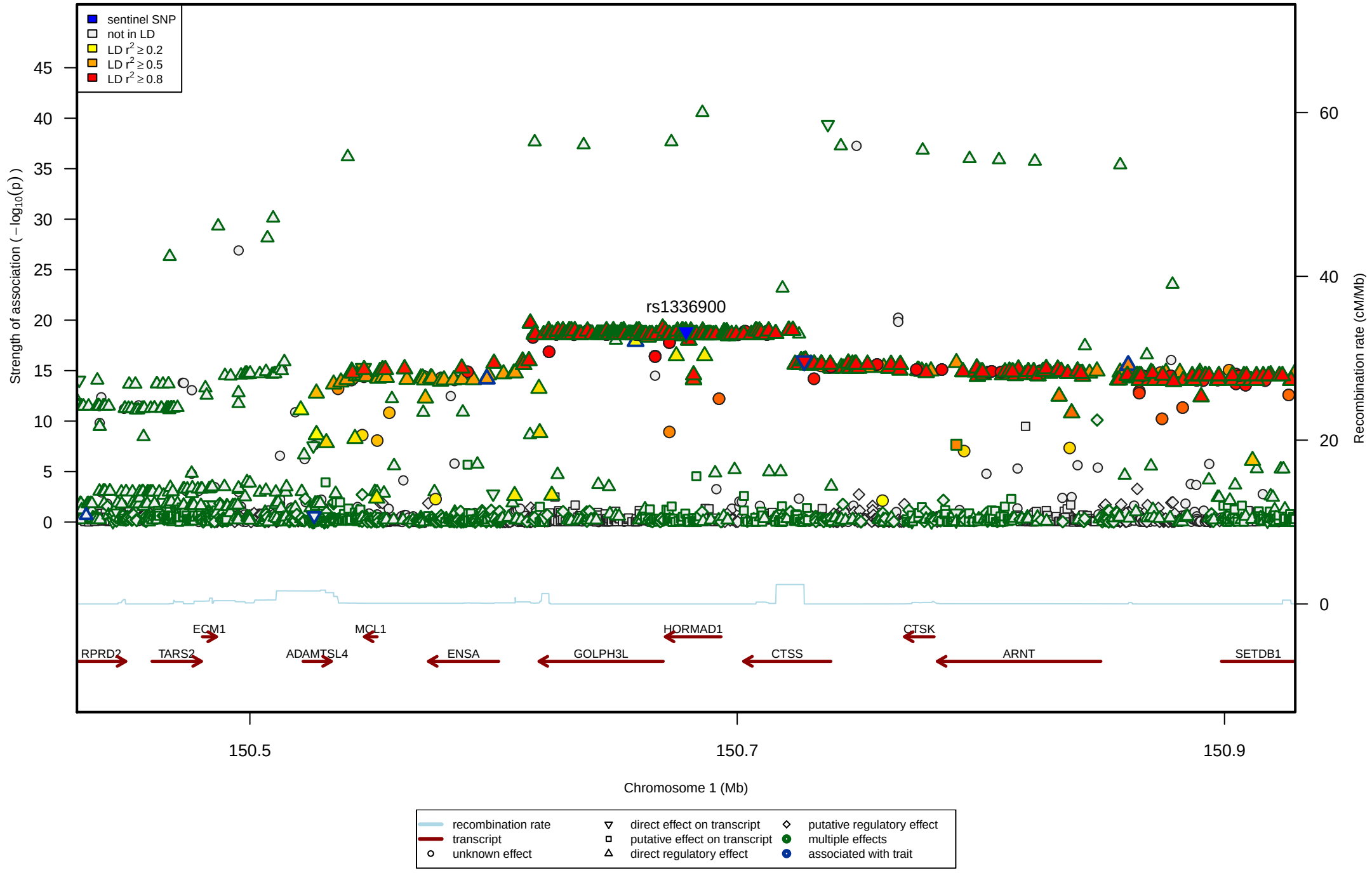

Locus: 201 Probe: 3181−50_2 Target gene(s): CTSS
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs1336900
Strength of association (−log10(p))
Recombination rate (cM/Mb)
ECM1
MCL1
HORMAD1
CTSK
RPRD2
TARS2
ADAMTSL4
ENSA
GOLPH3L
CTSS
ARNT
SETDB1
150.5
150.7
150.9
Chromosome 1 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait