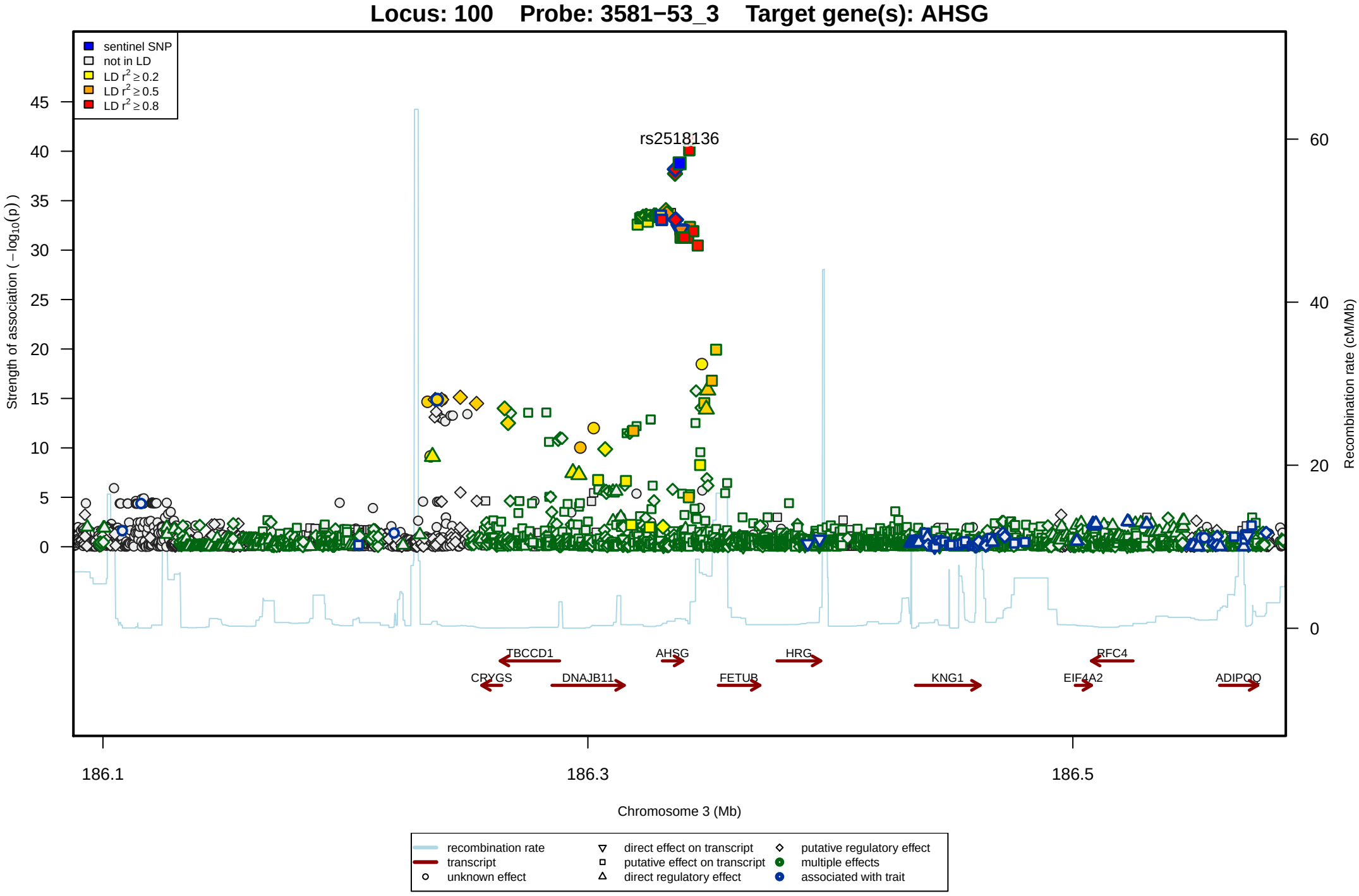

Locus: 100 Probe: 3581−53_3 Target gene(s): AHSG
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs2518136
Strength of association (−log10(p))
Recombination rate (cM/Mb)
TBCCD1
AHSG
HRG
RFC4
CRYGS
DNAJB11
FETUB
KNG1
EIF4A2
ADIPOQ
186.1
186.3
186.5
Chromosome 3 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait