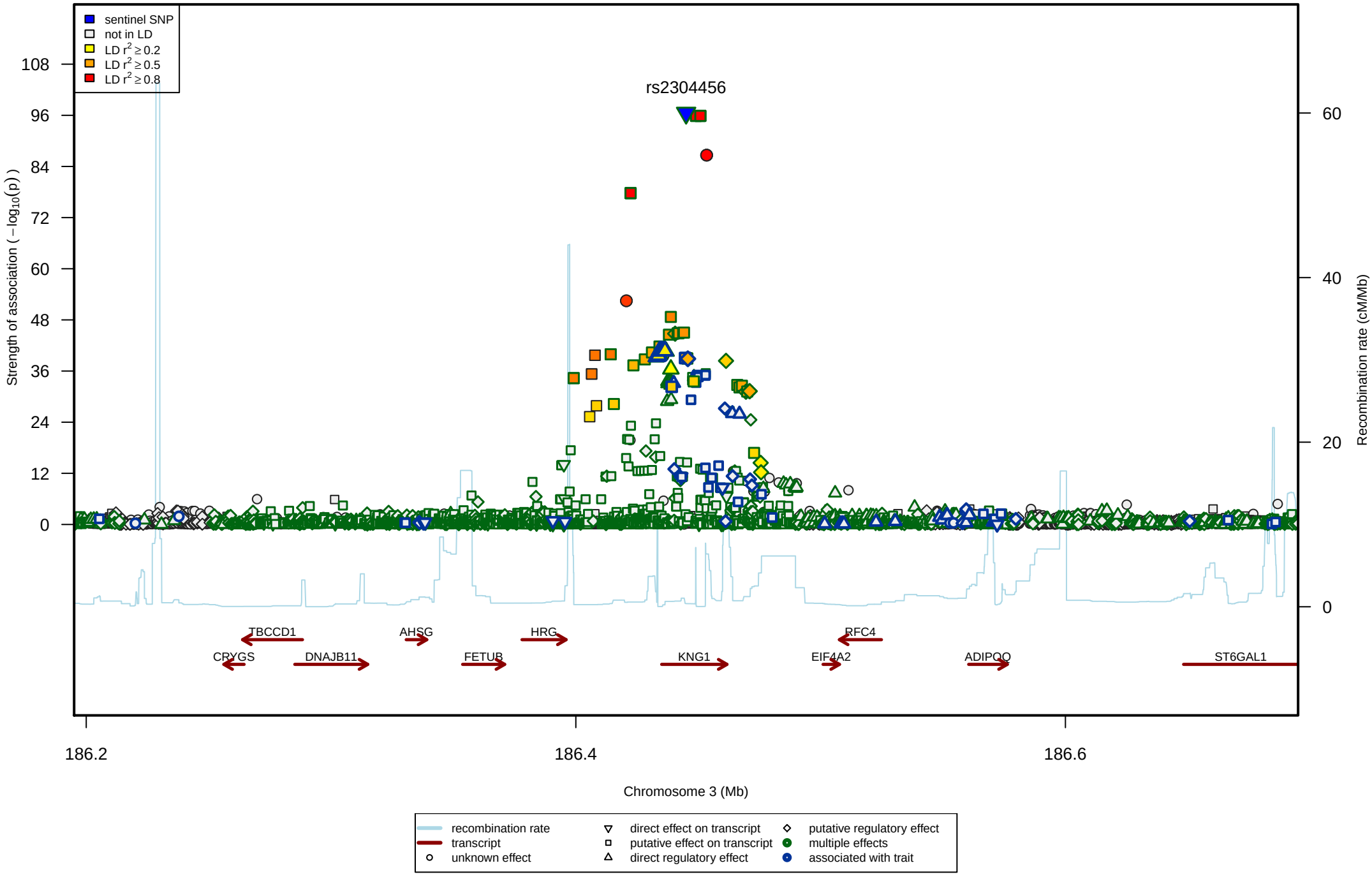

Locus: 26 Probe: 4495−33_2 Target gene(s): KNG1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs2304456
Strength of association (−log10(p))
Recombination rate (cM/Mb)
108
96
84
72
60
48
36
24
12
0
60
40
20
0
TBCCD1
AHSG
HRG
RFC4
CRYGS
DNAJB11
FETUB
KNG1
EIF4A2
ADIPOQ
ST6GAL1
186.2
186.4
186.6
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