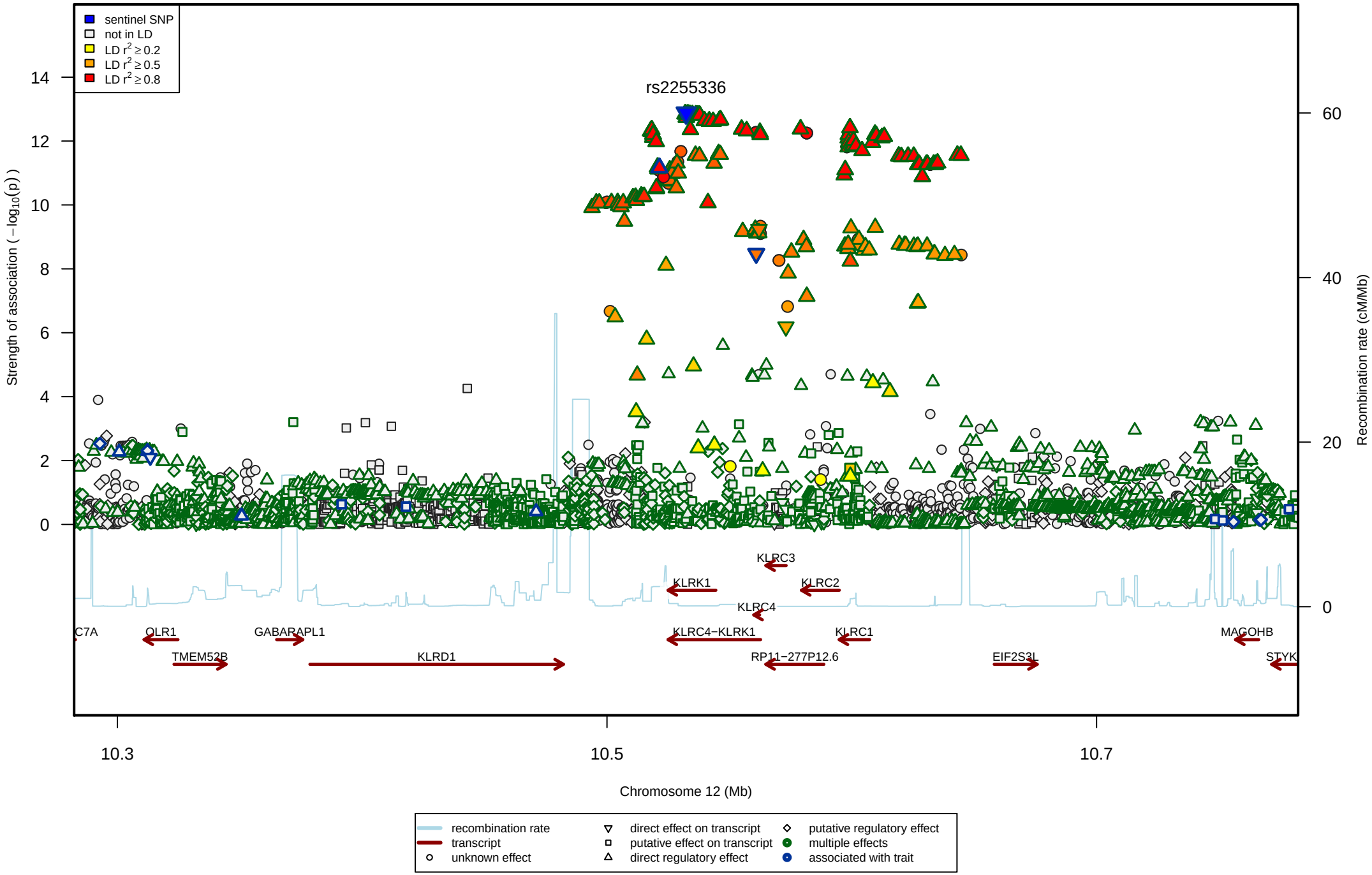

Locus: 332 Probe: 3056−11_1 Target gene(s): KLRK1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs2255336
Strength of association (−log10(p))
Recombination rate (cM/Mb)
KLRC3
KLRK1
KLRC2
KLRC4
C7A
OLR1
GABARAPL1
KLRC4−KLRK1
KLRC1
MAGOHB
TMEM52B
KLRD1
RP11−277P12.6
EIF2S3L
STYK
10.3
10.5
10.7
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