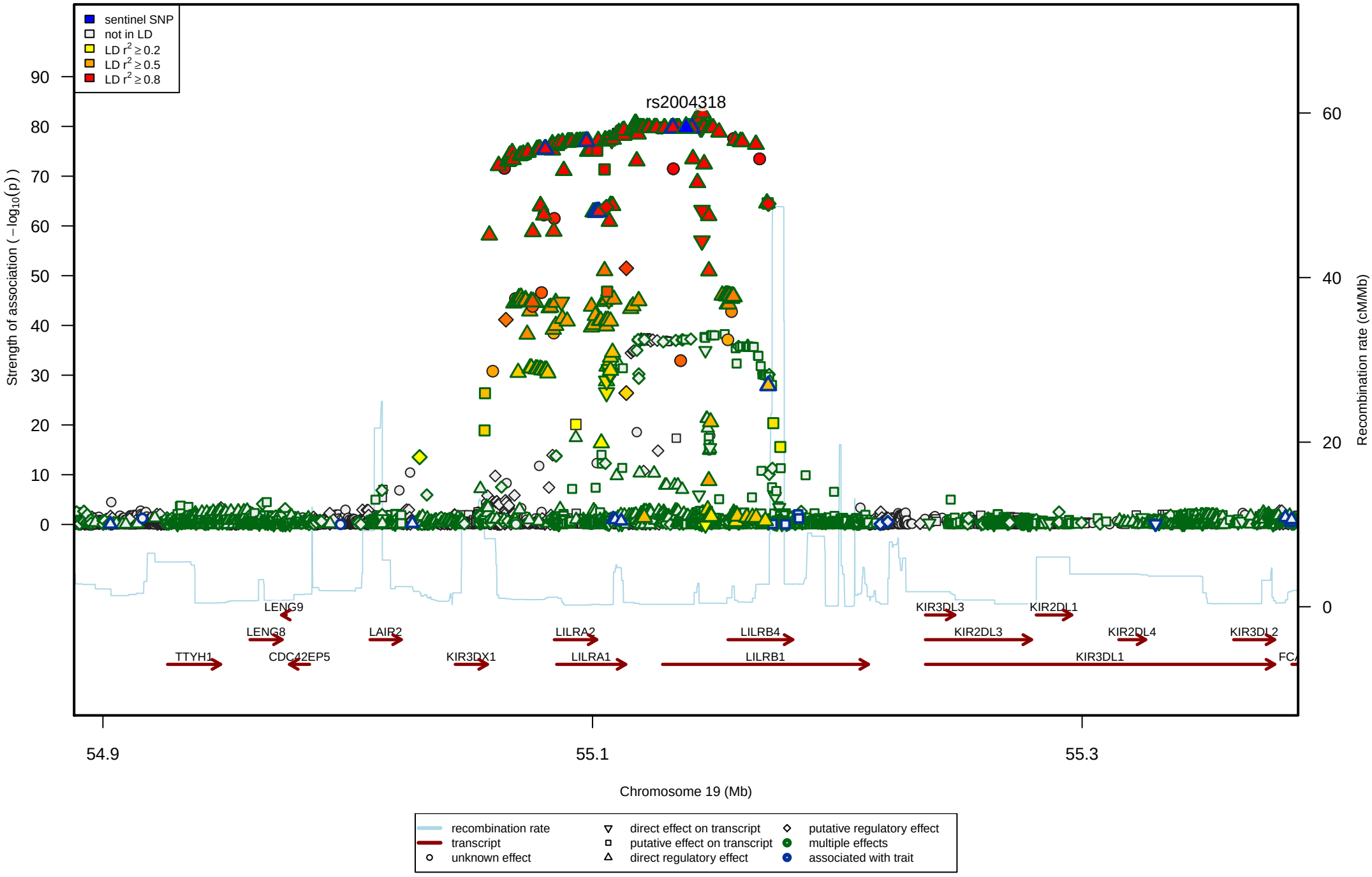

Locus: 41 Probe: 5090−49_2 Target gene(s): LILRB1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs2004318
Strength of association (−log10(p))
Recombination rate (cM/Mb)
LENG9
LENG8
LAIR2
LILRA2
LILRB4
KIR3DL3
KIR2DL1
KIR2DL3
KIR2DL4
KIR3DL2
TTYH1
CDC42EP5
KIR3DX1
LILRA1
LILRB1
KIR3DL1
54.9
55.1
55.3
Chromosome 19 (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