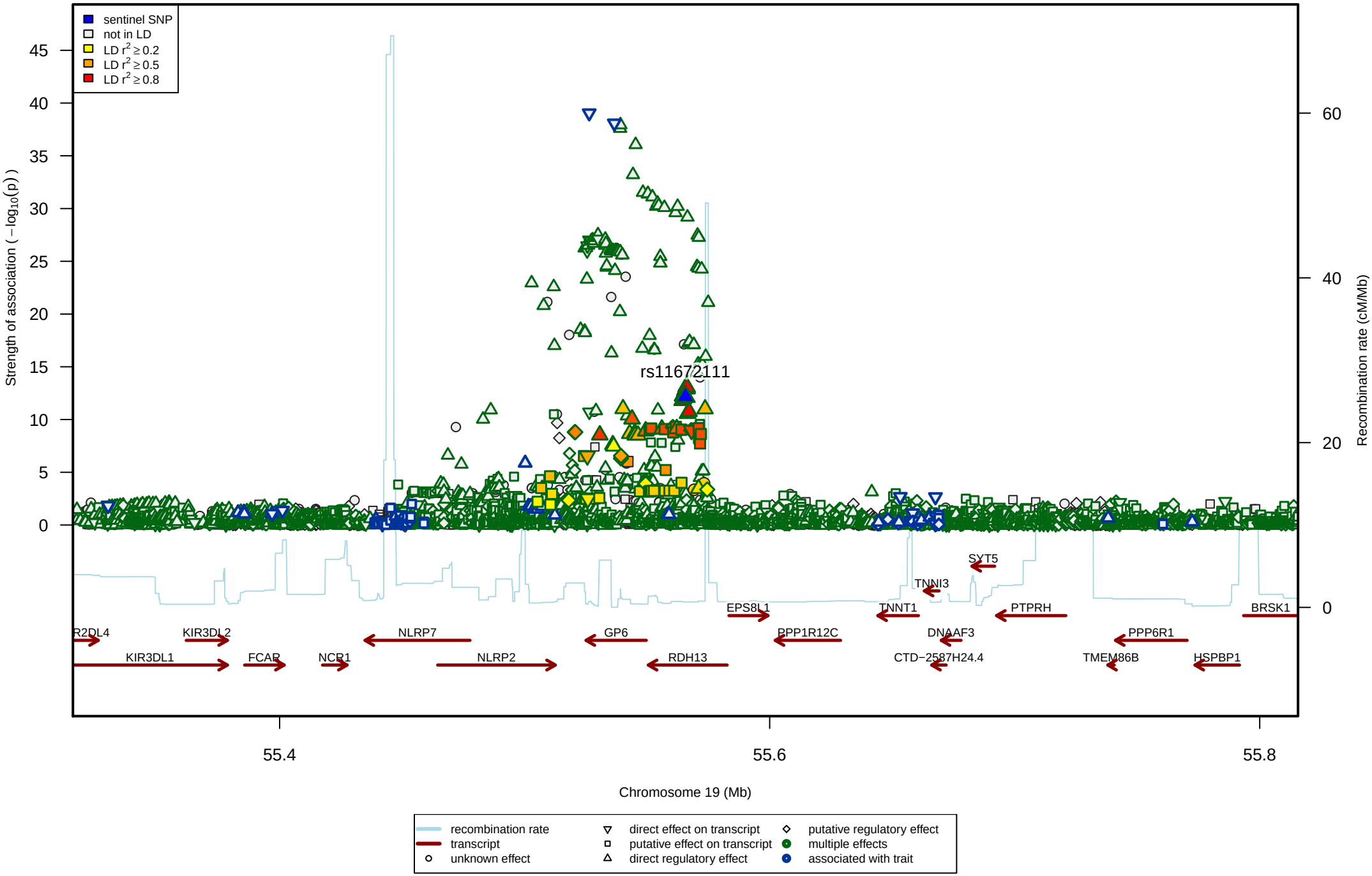

Locus: 355 Probe: 3194−36_2 Target gene(s): GP6
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
Strength of association (−log10(p))
Recombination rate (cM/Mb)
rs11672111
SYT5
TNNI3
EPS8L1
TNNT1
PTPRH
BRSK1
R2DL4
KIR3DL2
NLRP7
GP6
PPP1R12C
DNAAF3
PPP6R1
KIR3DL1
FCAR
NCR1
NLRP2
RDH13
CTD−2587H24.4
TMEM86B
HSPBP1
55.4
55.6
55.8
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