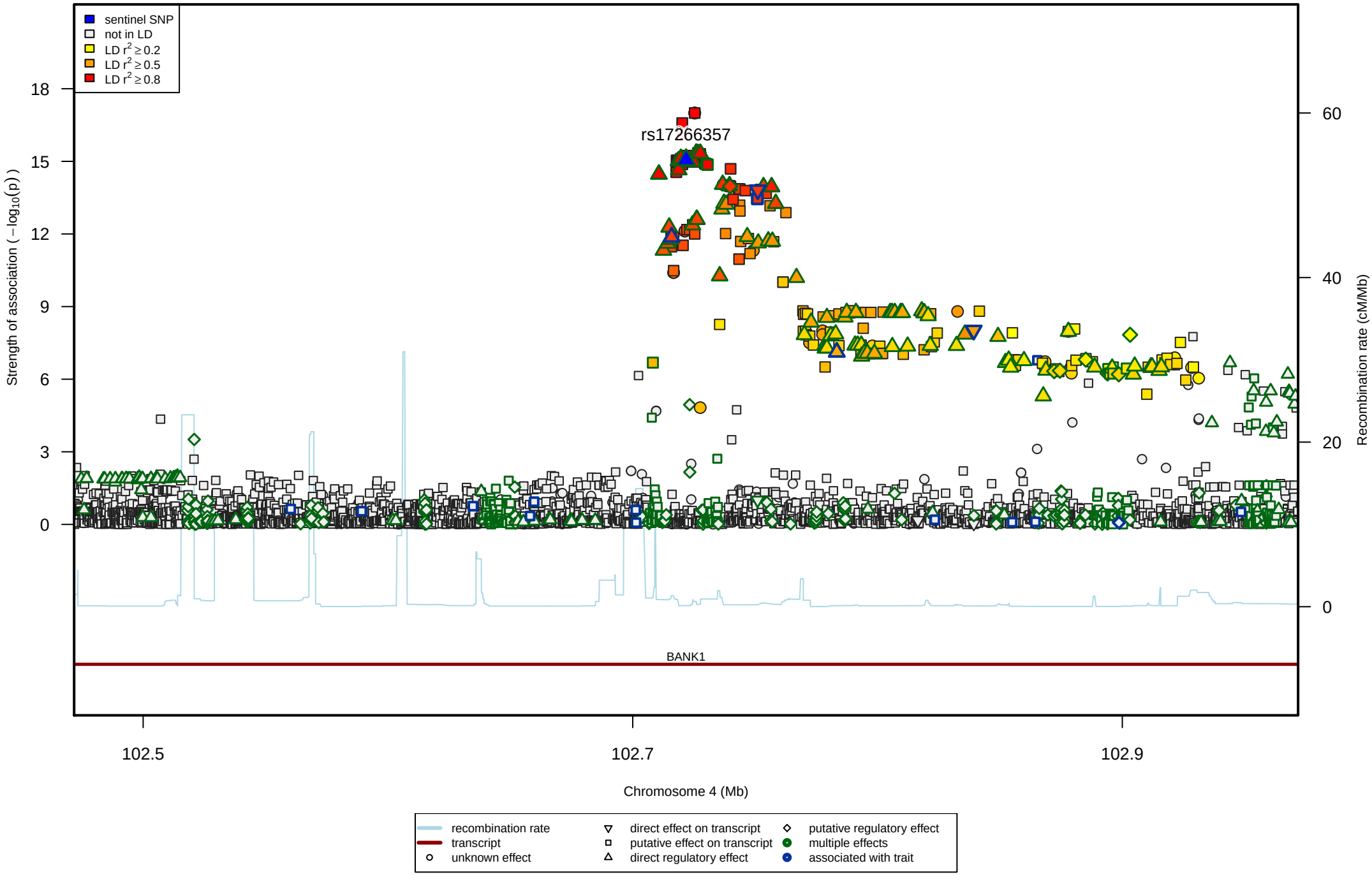

Locus: 262 Probe: 4903−72_1 Target gene(s): PPP3CA PPP3R1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs17266357
BANK1
Strength of association (−log10(p))
Recombination rate (cM/Mb)
Chromosome 4 (Mb)
102.5
102.7
102.9
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait