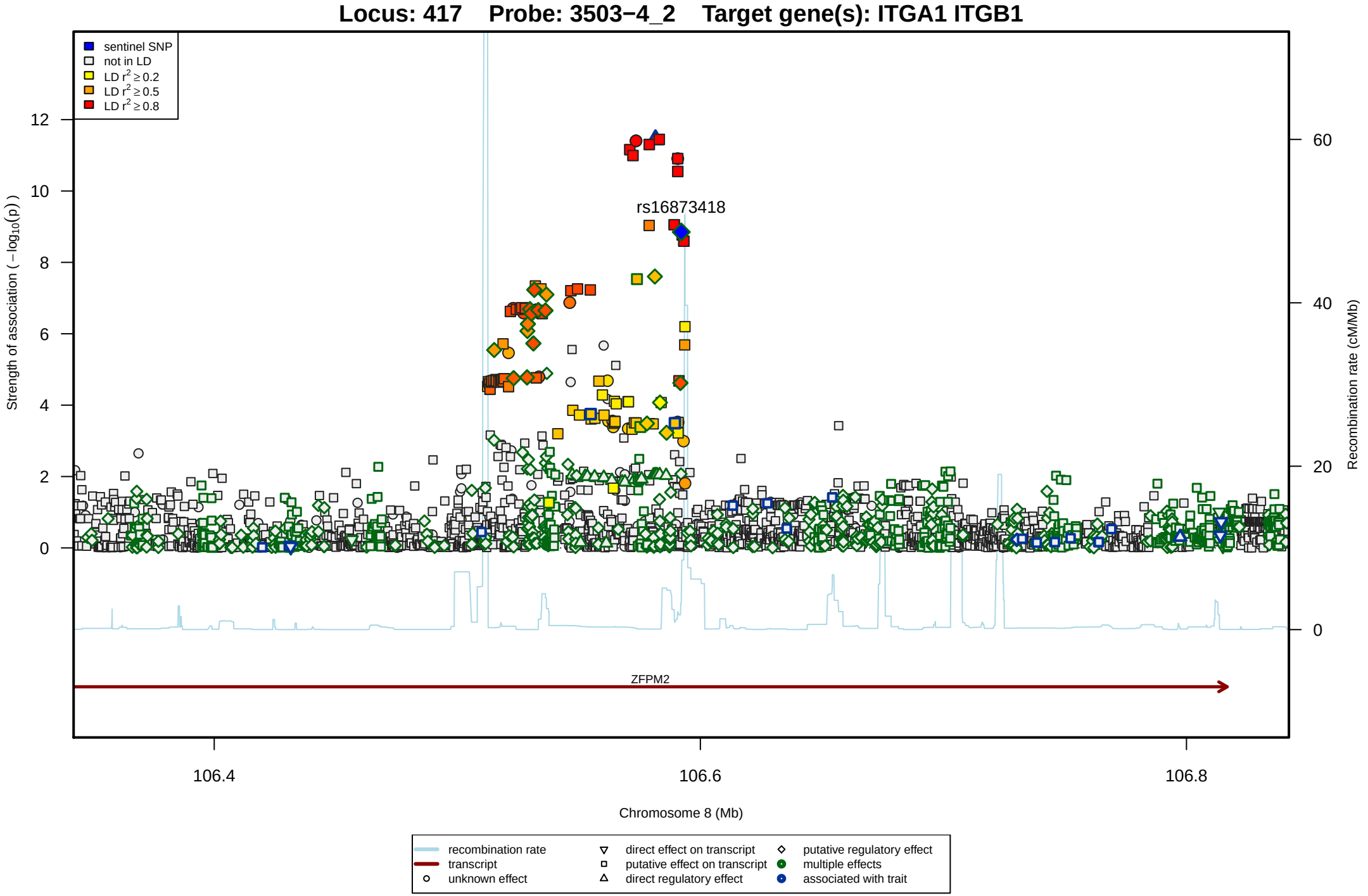

Locus: 417 Probe: 3503−4_2 Target gene(s): ITGA1 ITGB1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs16873418
ZFPM2
Strength of association (−log10(p))
Recombination rate (cM/Mb)
Chromosome 8 (Mb)
106.4
106.6
106.8
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait