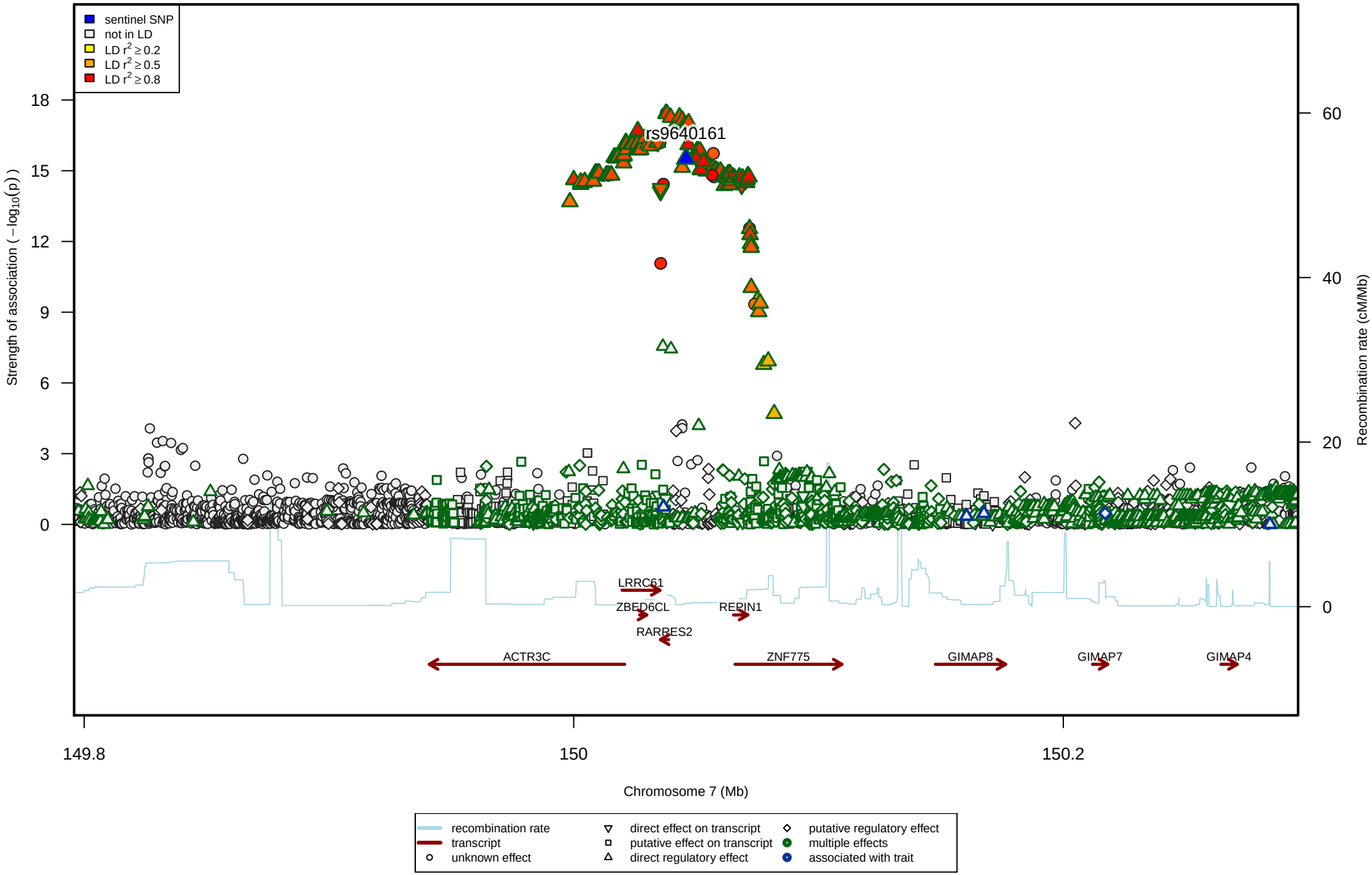

Locus: 249 Probe: 3079−62_2 Target gene(s): RARRES2
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs9640161
Strength of association (−log10(p))
Recombination rate (cM/Mb)
LRRC61
ZBED6CL
REPIN1
RARRES2
ACTR3C
ZNF775
GIMAP8
GIMAP7
GIMAP4
149.8
150
150.2
Chromosome 7 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait