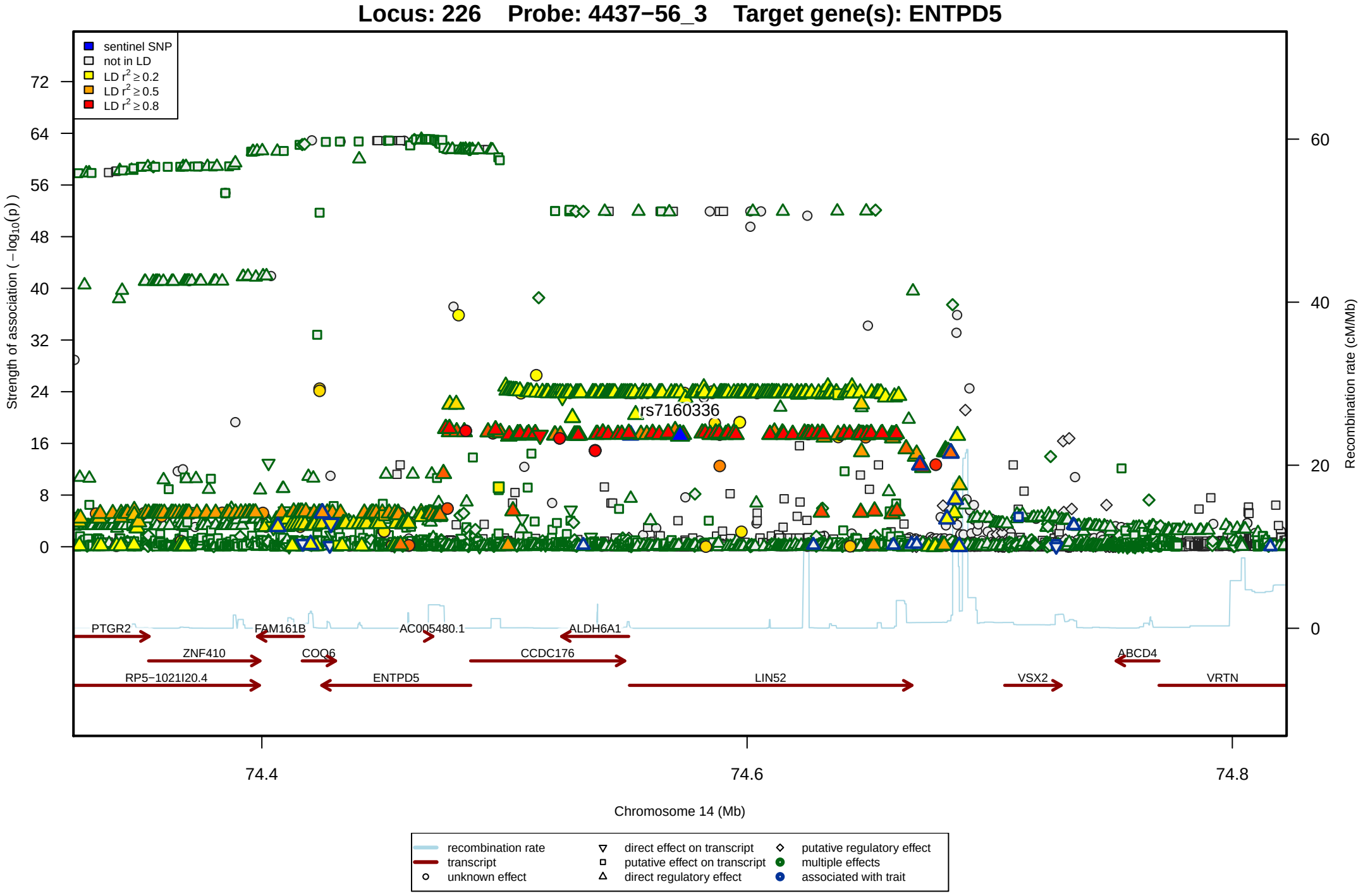

Locus: 226 Probe: 4437−56_3 Target gene(s): ENTPD5
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
Strength of association (−log10(p))
Recombination rate (cM/Mb)
rs7160336
PTGR2
FAM161B
AC005480.1
ALDH6A1
ZNF410
COQ6
CCDC176
ABCD4
RP5−1021I20.4
ENTPD5
LIN52
VSX2
VRTN
74.4
74.6
74.8
Chromosome 14 (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