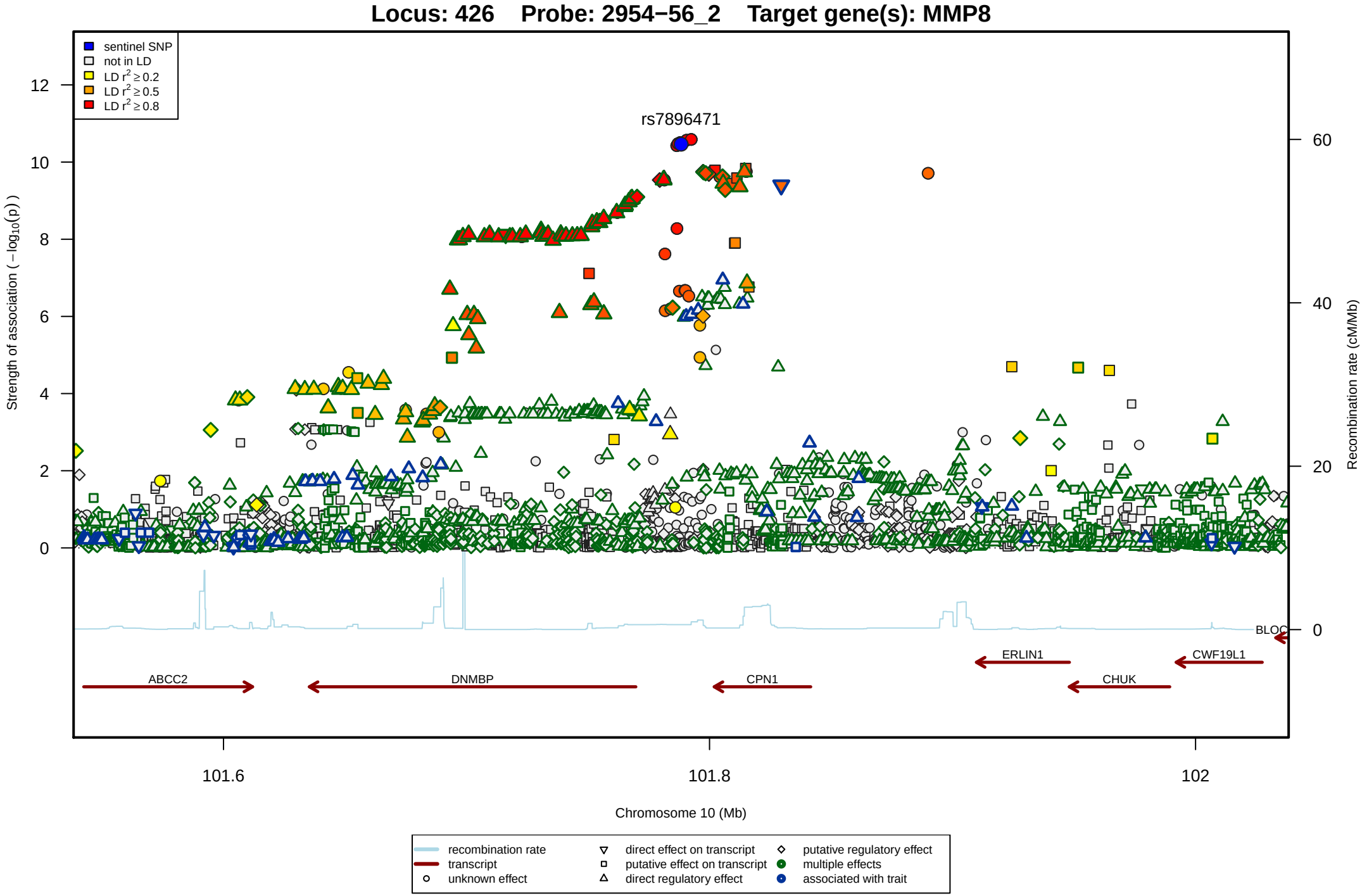

Locus: 426 Probe: 2954−56_2 Target gene(s): MMP8
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs7896471
Strength of association (−log10(p))
Recombination rate (cM/Mb)
ABCC2
DNMBP
CPN1
ERLIN1
CHUK
CWF19L1
BLOC
101.6
101.8
102
Chromosome 10 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait