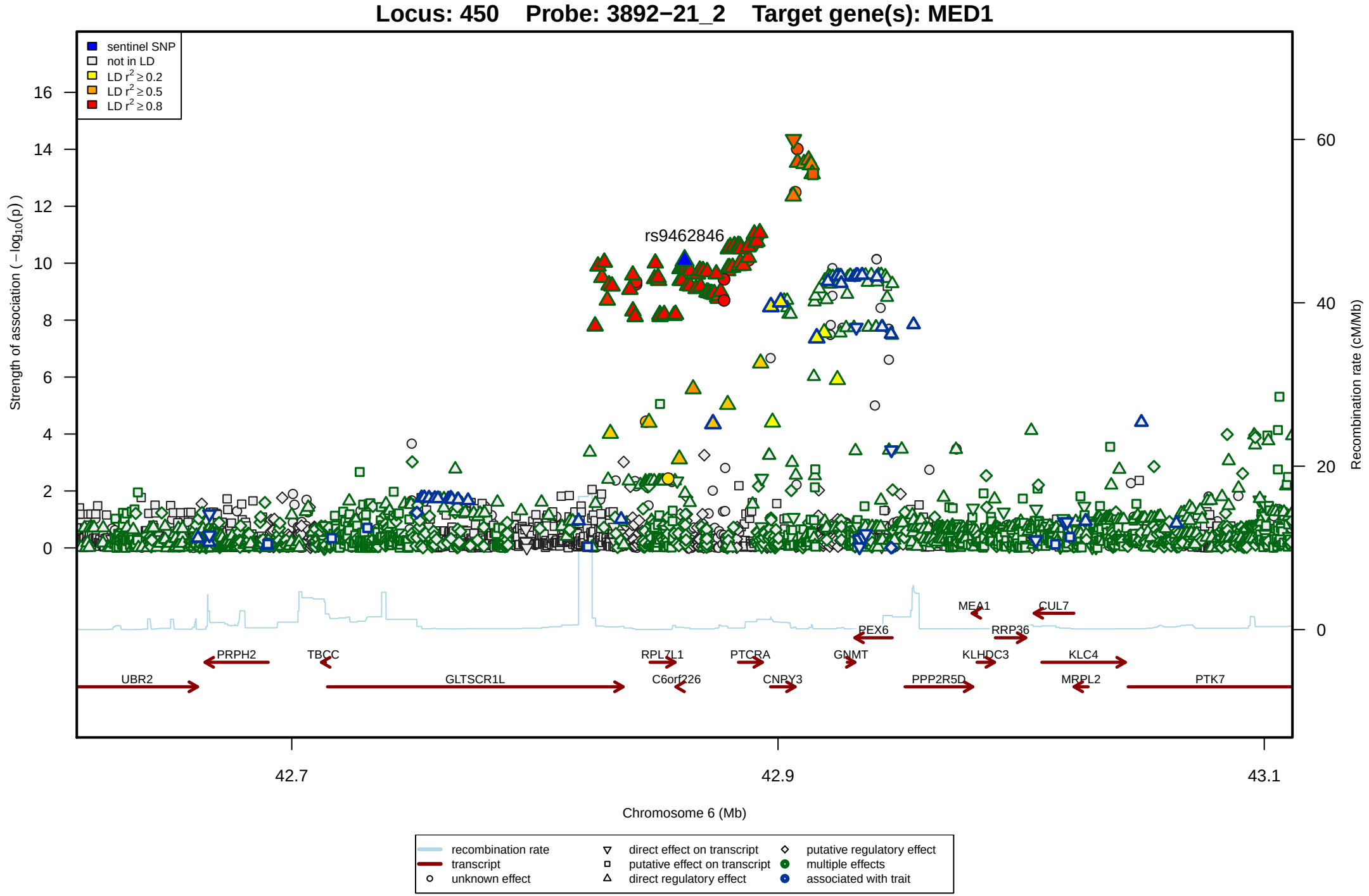

Locus: 450 Probe: 3892−21_2 Target gene(s): MED1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs9462846
Strength of association (−log10(p))
Recombination rate (cM/Mb)
Chromosome 6 (Mb)
42.7
42.9
43.1
MEA1
CUL7
PEX6
RRP36
PRPH2
TBCC
RPL7L1
PTCRA
GNMT
KLHDC3
KLC4
UBR2
GLTSCR1L
C6orf226
CNPY3
PPP2R5D
MRPL2
PTK7
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait