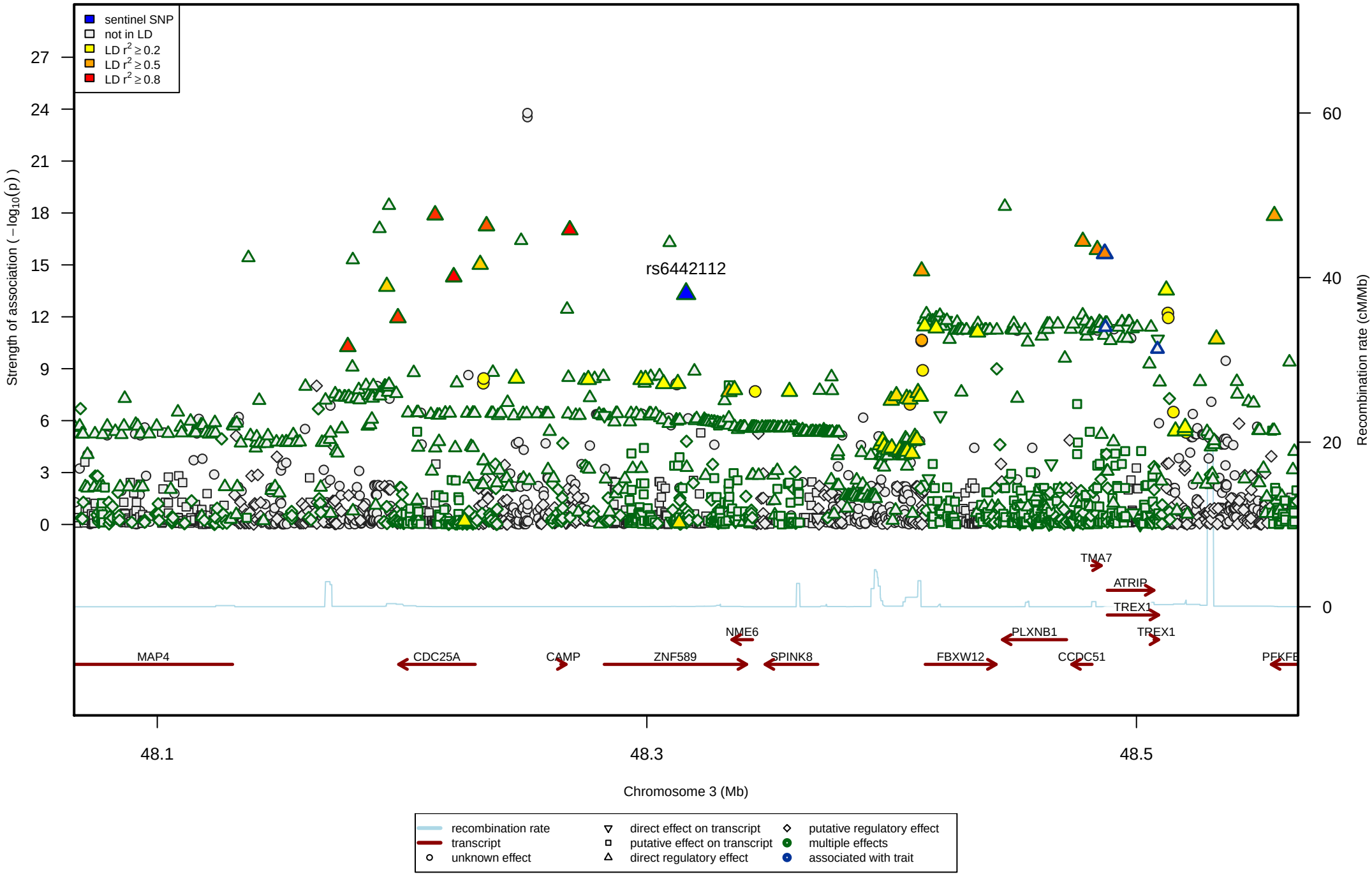

Locus: 313 Probe: 4407−10_1 Target gene(s): MST1
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs6442112
Strength of association (−log10(p))
Recombination rate (cM/Mb)
Chromosome 3 (Mb)
48.1
48.3
48.5
MAP4
CDC25A
CAMP
ZNF589
SPINK8
NME6
FBXW12
PLXNB1
CCDC51
TREX1
ATRIP
TMA7
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait