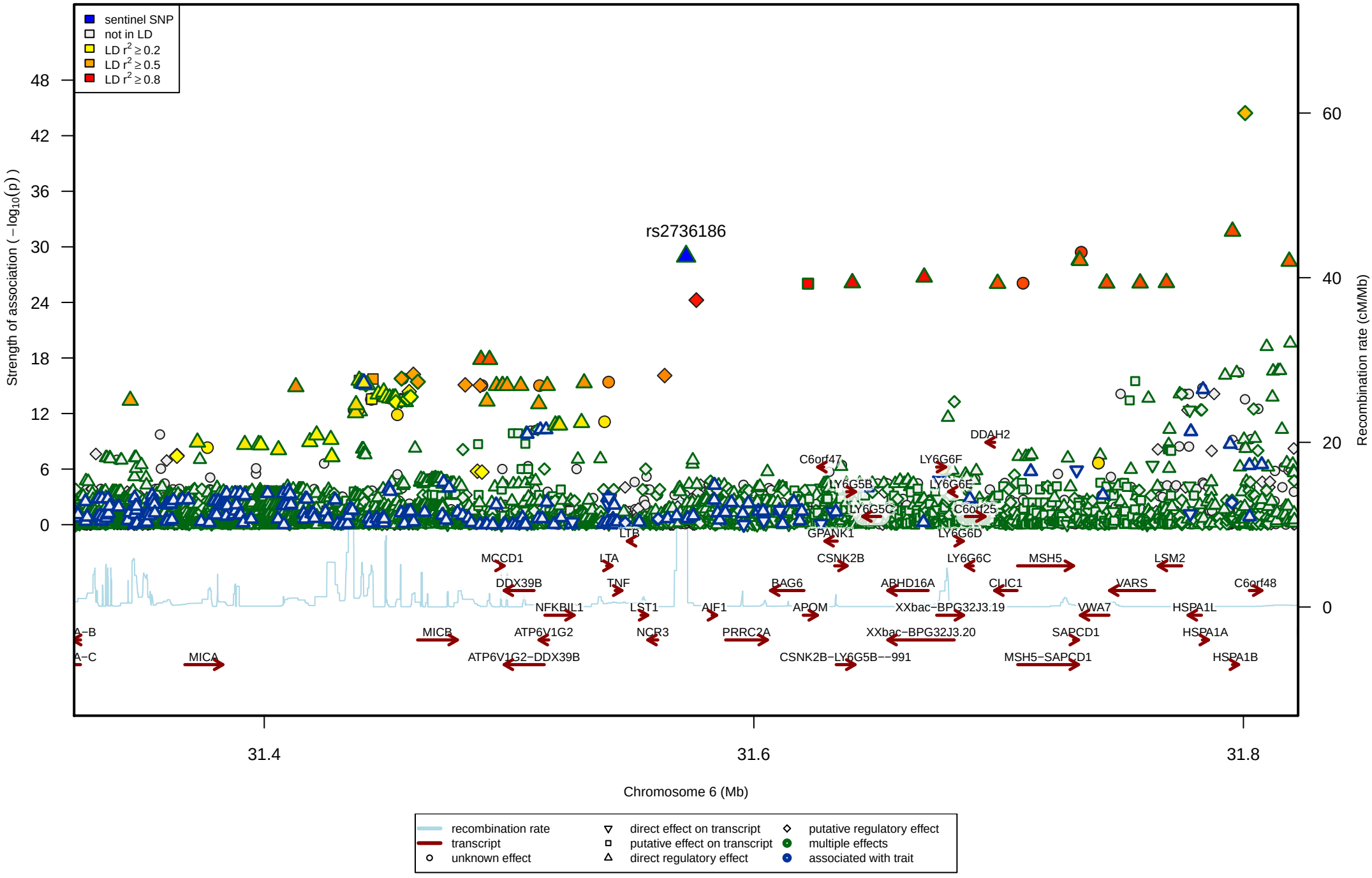

Locus: 132 Probe: 2954−56_2 Target gene(s): MMP8
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
Strength of association (−log10(p))
Recombination rate (cM/Mb)
rs2736186
DDAH2
C6orf47
LY6G6F
LY6G5B
LY6G6E
LY6G5C
C6orf25
GPANK1
LY6G6D
LTB
MCCD1
LTA
CSNK2B
LY6G6C
MSH5
LSM2
DDX39B
TNF
BAG6
ABHD16A
CLIC1
VARS
C6orf48
NFKBIL1
LST1
AIF1
APOM
XXbac−BPG32J3.19
VWA7
HSPA1L
A−B
MICB
ATP6V1G2
NCR3
PRRC2A
XXbac−BPG32J3.20
SAPCD1
HSPA1A
A−C
MICA
ATP6V1G2−DDX39B
CSNK2B−LY6G5B−−991
MSH5−SAPCD1
HSPA1B
31.4
31.6
31.8
Chromosome 6 (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