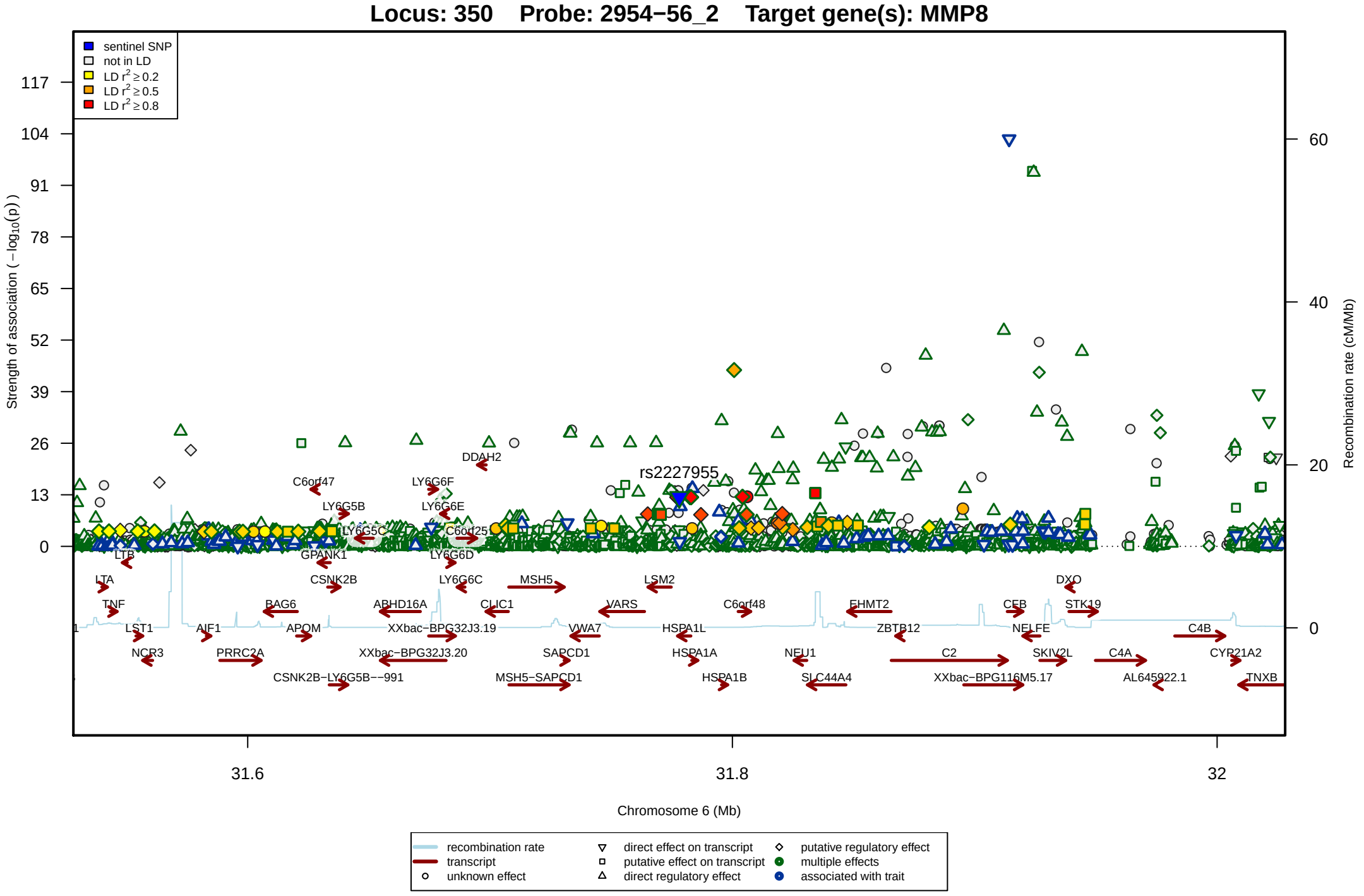

Locus: 350 Probe: 2954−56_2 Target gene(s): MMP8
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
Strength of association (−log10(p))
Recombination rate (cM/Mb)
rs2227955
DDAH2
C6orf47
LY6G6F
LY6G5B
LY6G6E
LY6G5C
C6orf25
GPANK1
LY6G6D
LTB
LTA
CSNK2B
LY6G6C
MSH5
LSM2
DXO
TNF
BAG6
APHD16A
CLIC1
VARS
C6orf48
EHMT2
CEB
STK19
LST1
AIF1
APOM
XXbac−BPG32J3.19
VWA7
HSPA1L
ZBTB12
NELFE
C4B
NCR3
PRRC2A
XXbac−BPG32J3.20
SAPCD1
HSPA1A
NEU1
C2
SKIV2L
C4A
CYP21A2
CSNK2B−LY6G5B−−991
MSH5−SAPCD1
HSPA1B
SLC44A4
XXbac−BPG116M5.17
AL645922.1
TNXB
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