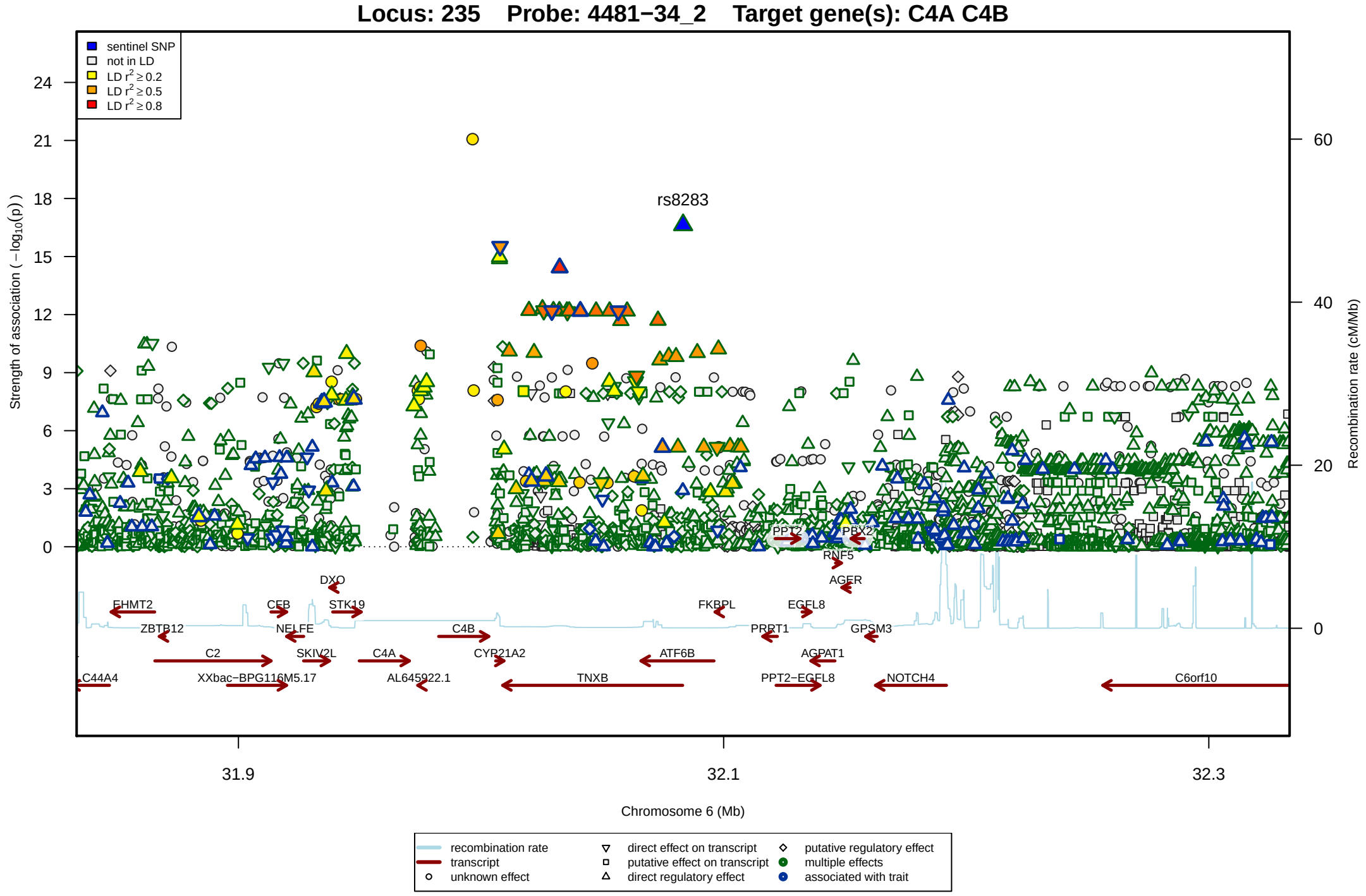

Locus: 235 Probe: 4481−34_2 Target gene(s): C4A C4B
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs8283
Strength of association (−log10(p))
Recombination rate (cM/Mb)
Chromosome 6 (Mb)
31.9
32.1
32.3
EHMT2
CFB
STK19
DXO
ZBTB12
NELFE
C4B
FKBPL
EGFL8
PRRT1
GPSM3
C2
SKIV2L
C4A
CYP21A2
ATF6B
AGPAT1
C44A4
XXbac−BPG116M5.17
AL645922.1
TNXB
PPT2−EGFL8
NOTCH4
C6orf10
RNF5
AGER
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait