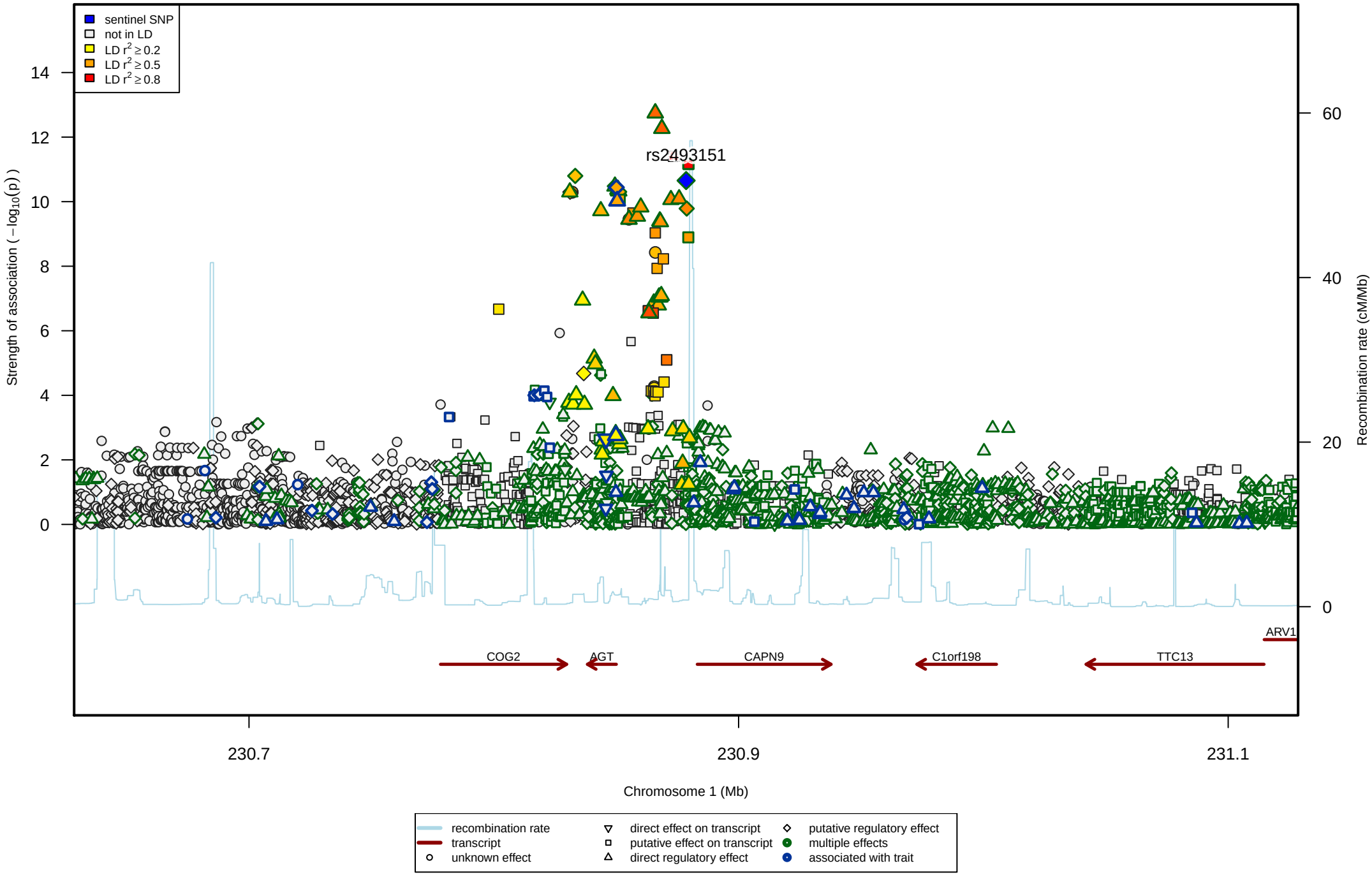

Locus: 420 Probe: 3484−60_2 Target gene(s): AGT
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs2493151
Strength of association (−log10(p))
Recombination rate (cM/Mb)
COG2
AGT
CAPN9
C1orf198
TTC13
ARV1
230.7
230.9
231.1
Chromosome 1 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait