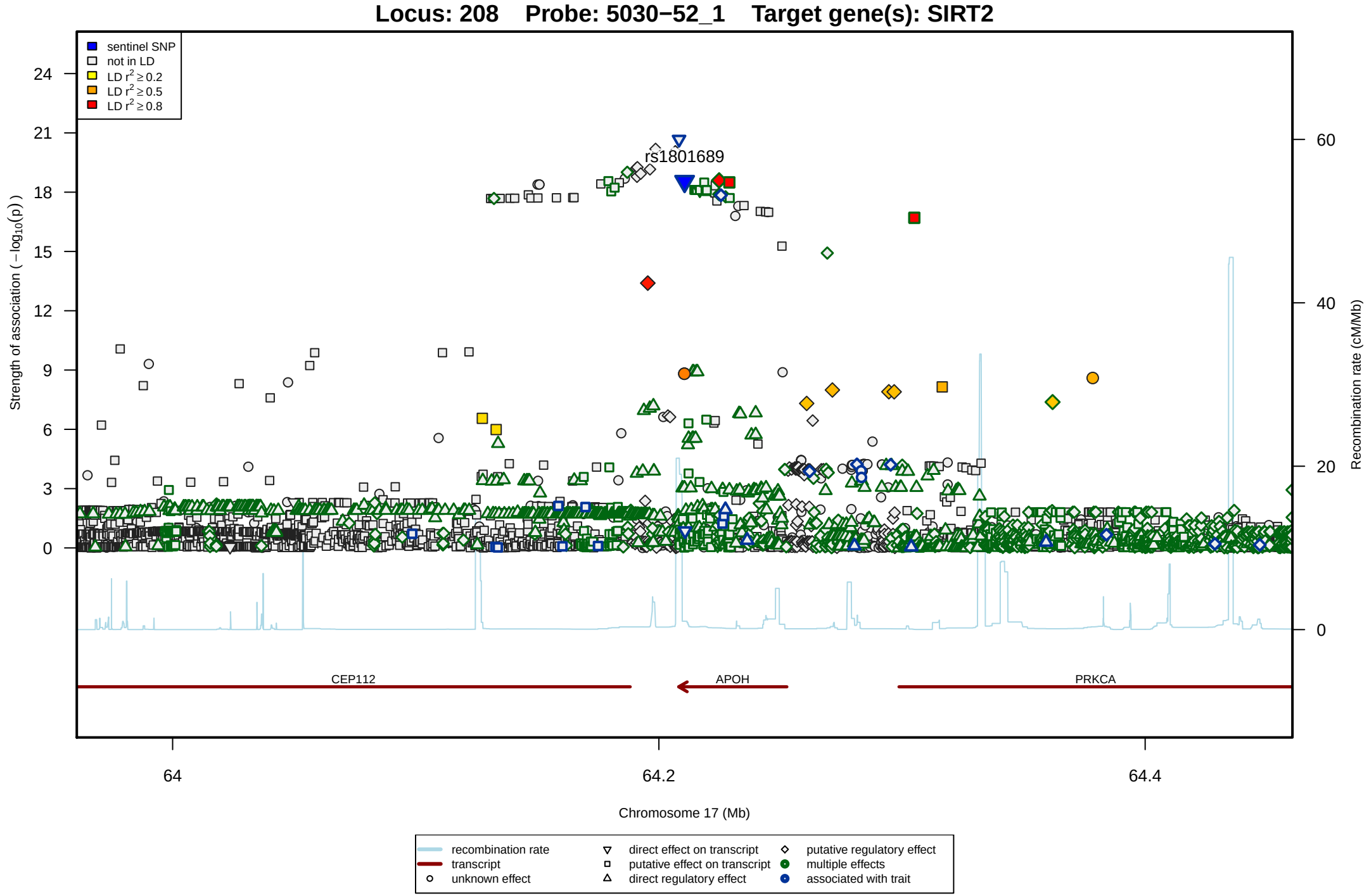

Locus: 208 Probe: 5030−52_1 Target gene(s): SIRT2
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs1801689
Strength of association (−log10(p))
Recombination rate (cM/Mb)
CEP112
APOH
PRKCA
Chromosome 17 (Mb)
recombination rate
transcript
unknown effect
direct effect on transcript
putative effect on transcript
direct regulatory effect
putative regulatory effect
multiple effects
associated with trait