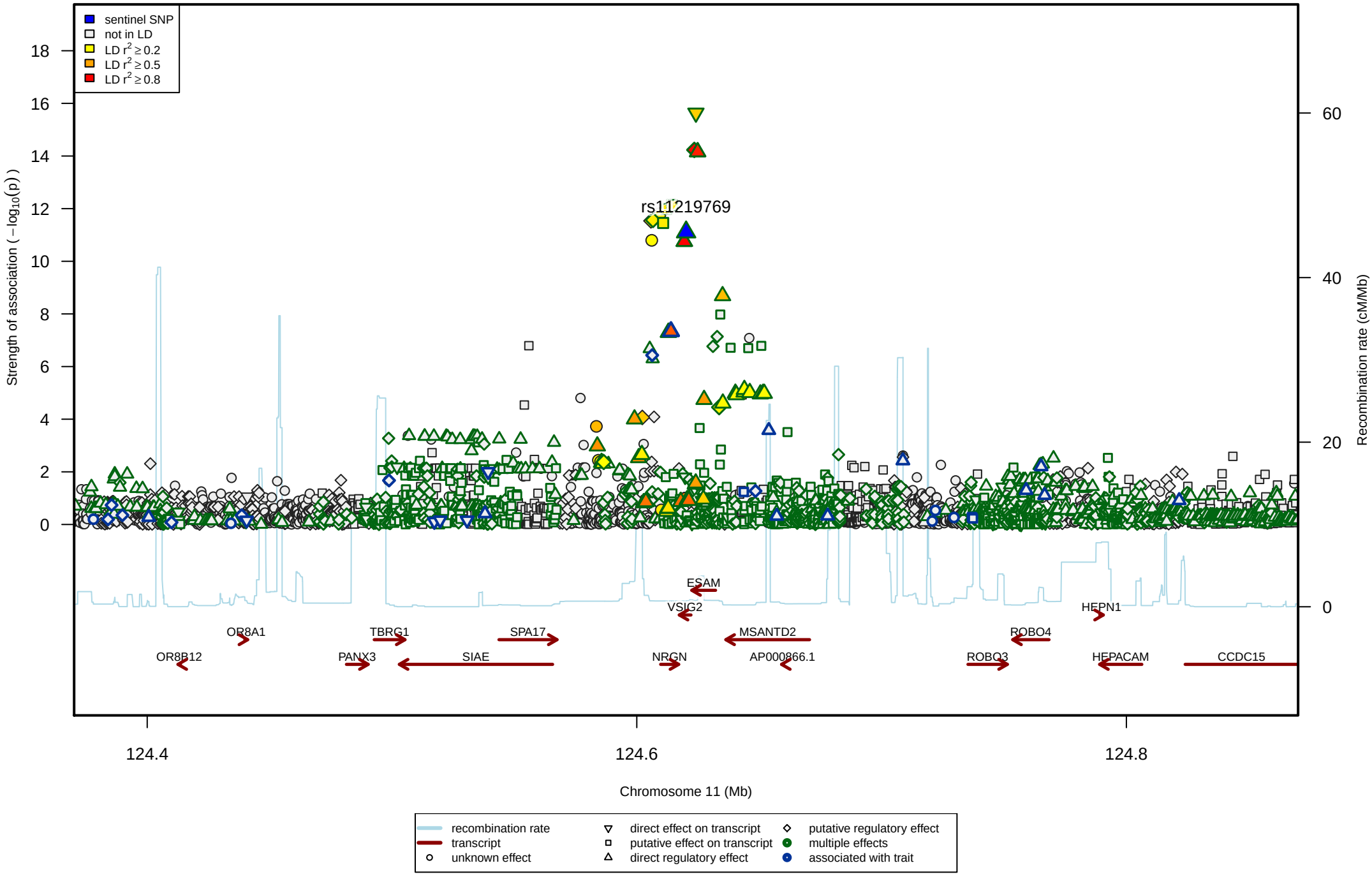

Locus: 406 Probe: 2981−9_3 Target gene(s): ESAM
sentinel SNP
not in LD
LD r2 ≥ 0.2
LD r2 ≥ 0.5
LD r2 ≥ 0.8
rs11219769
Strength of association (−log10(p))
Recombination rate (cM/Mb)
ESAM
VSIG2
HEPN1
OR8A1
TBRG1
SPA17
MSANTD2
ROBO4
OR8B12
PANX3
SIAE
NRGN
AP000866.1
ROBO3
HEPACAM
CCDC15
124.4
124.6
124.8
Chromosome 11 (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