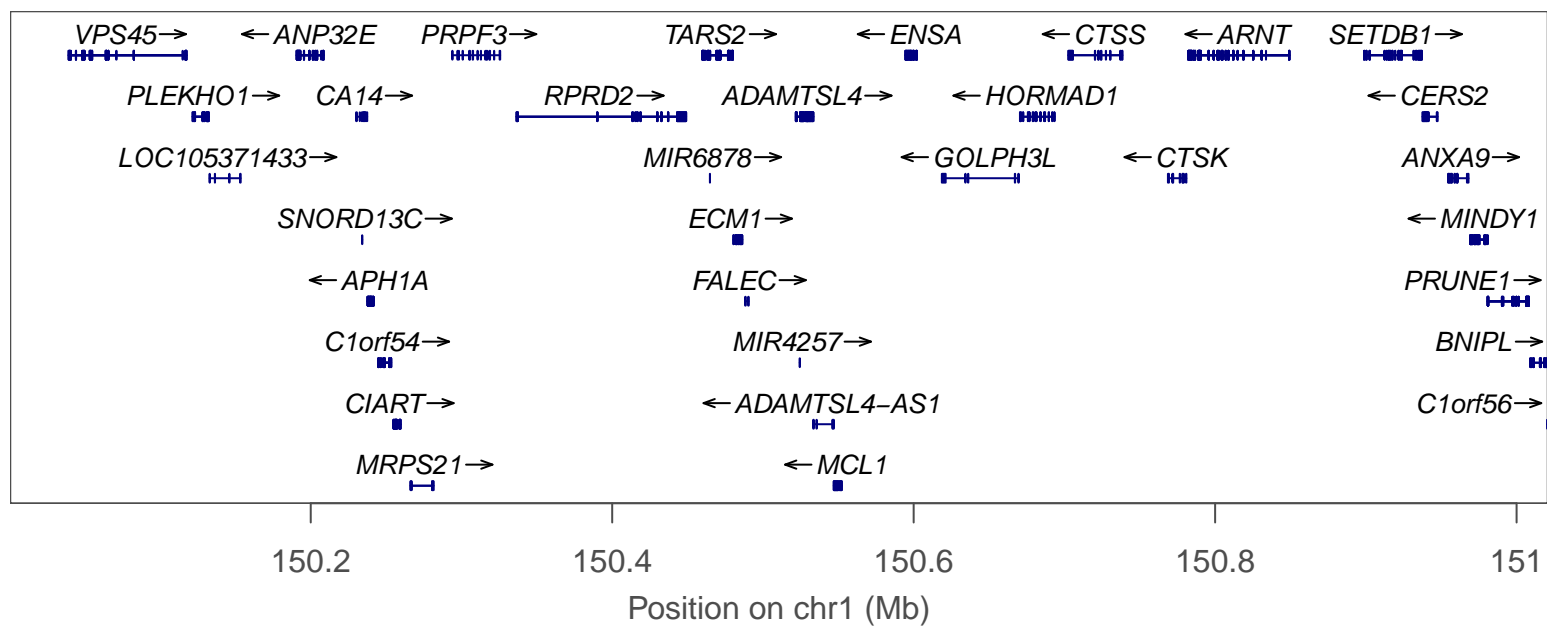
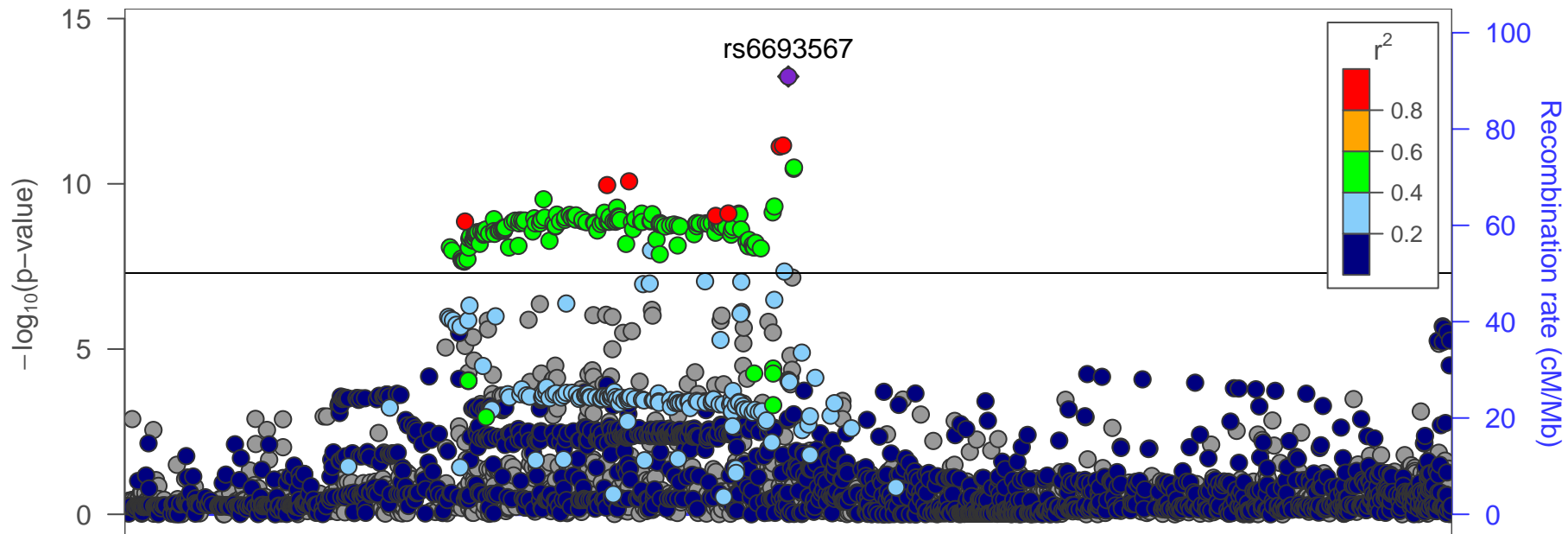
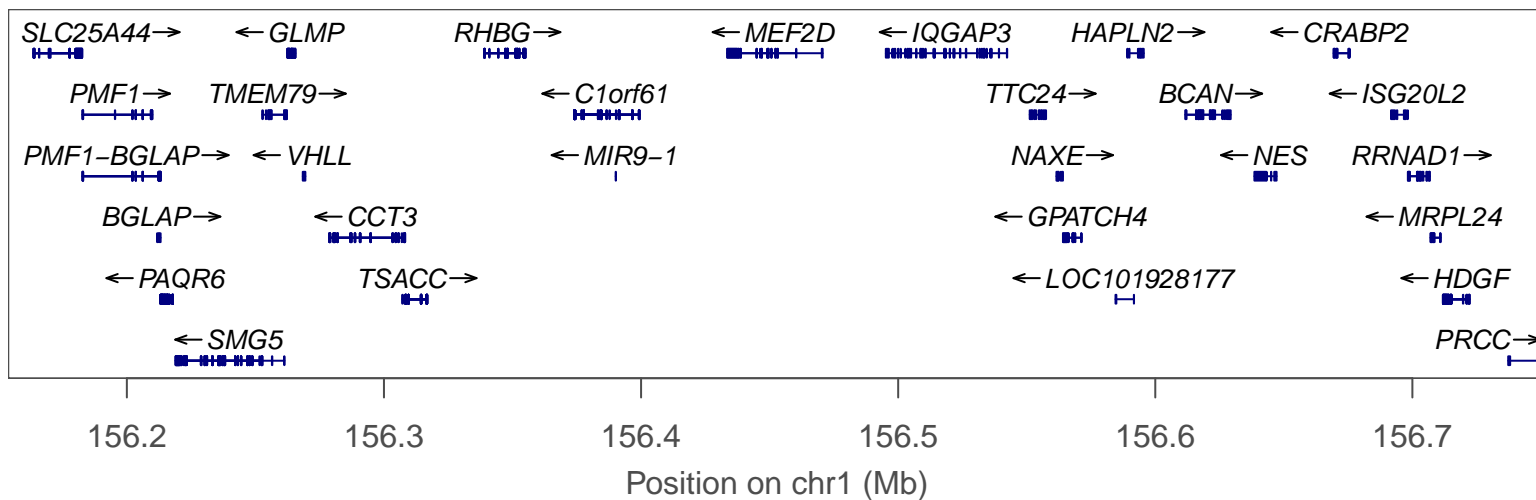
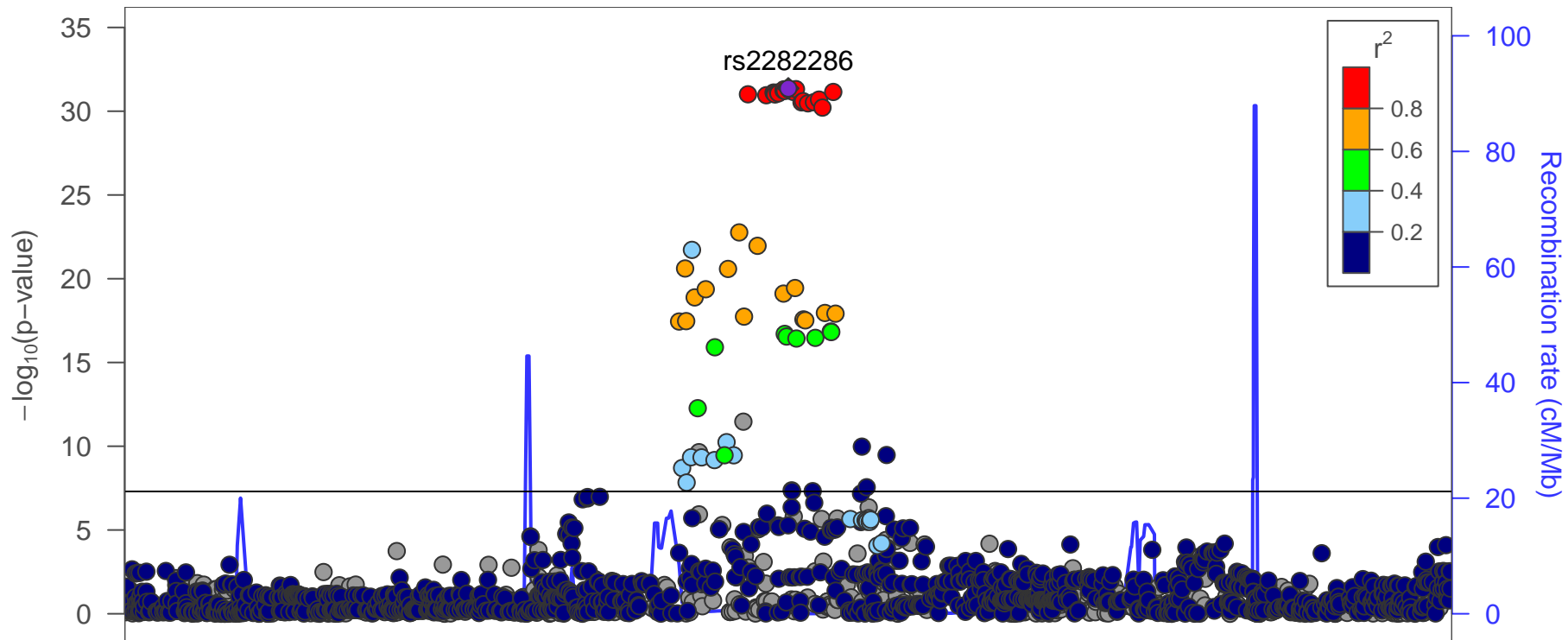


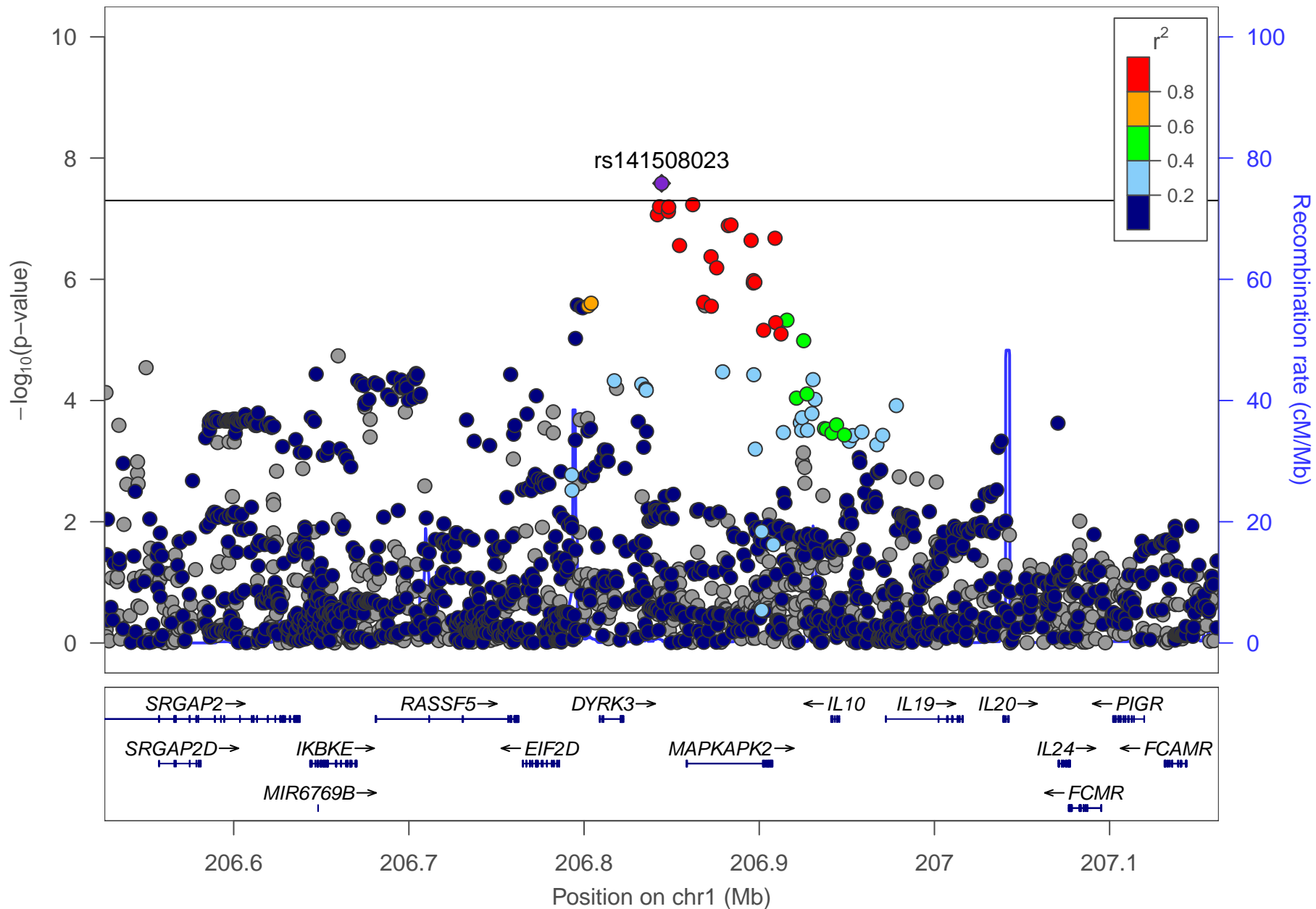
1 gene omitted

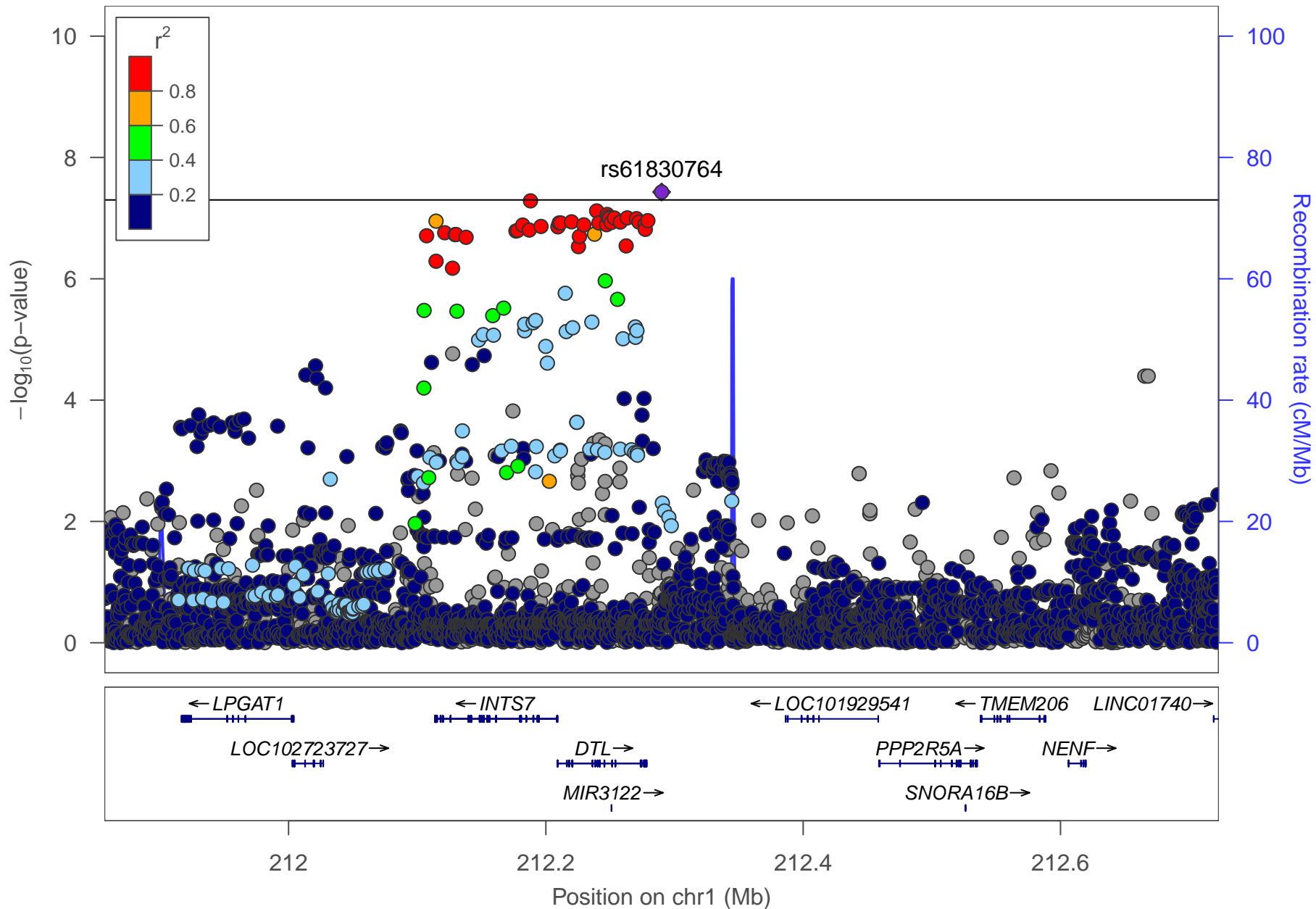
149.7                      149.8                      149.9                      150                      150.1

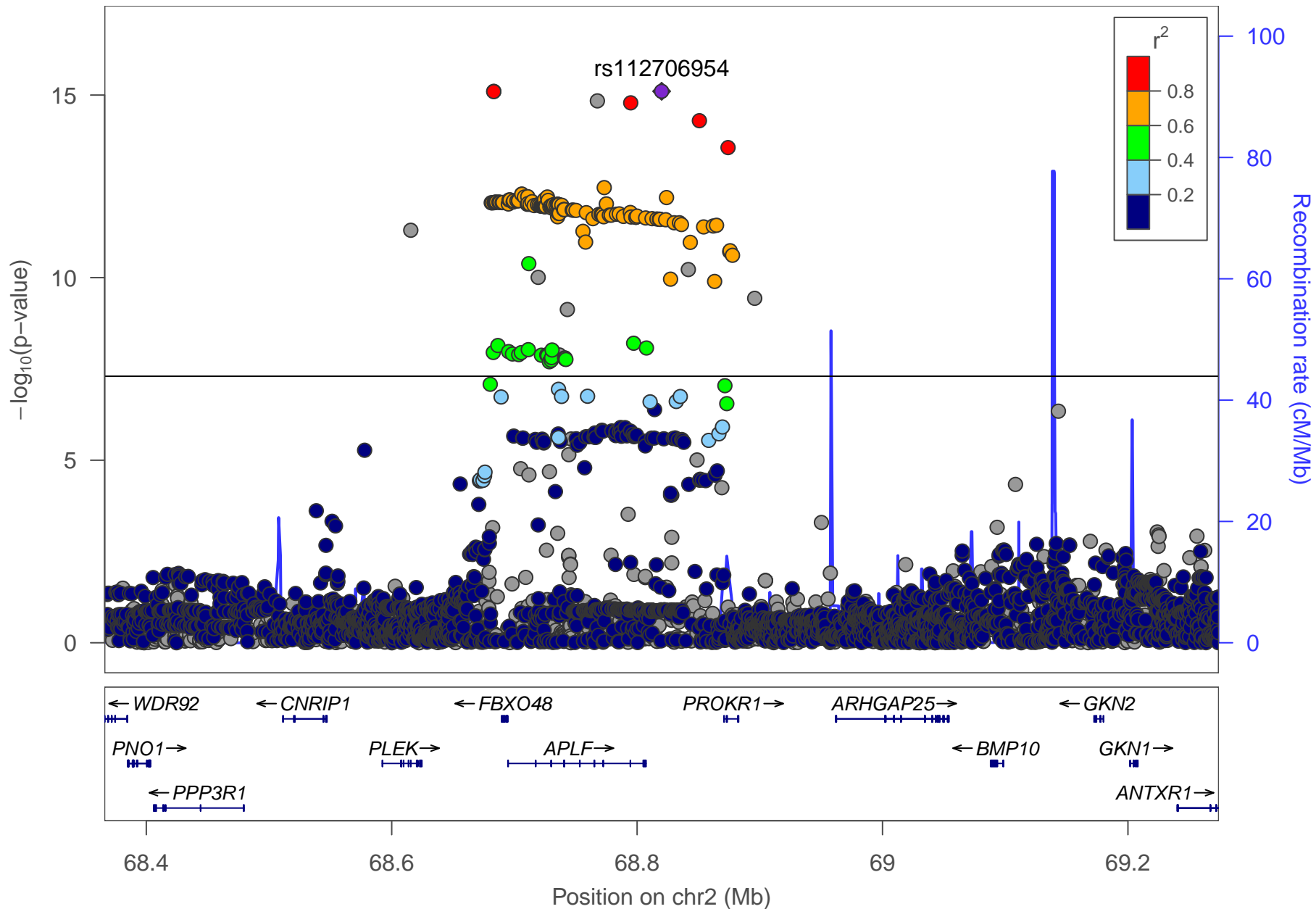
Position on chr1 (Mb)

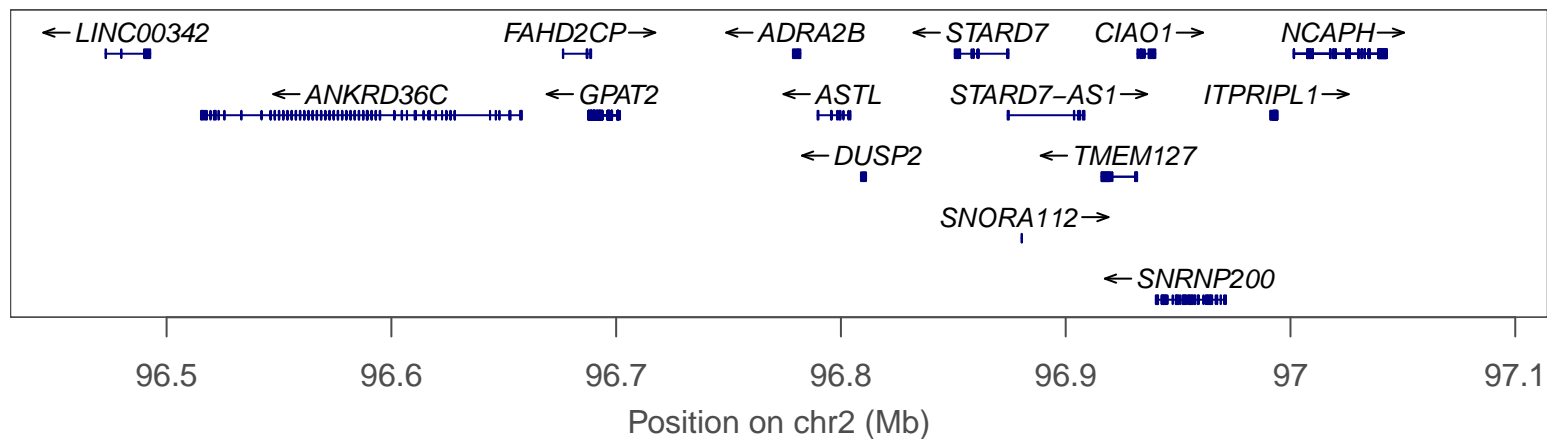
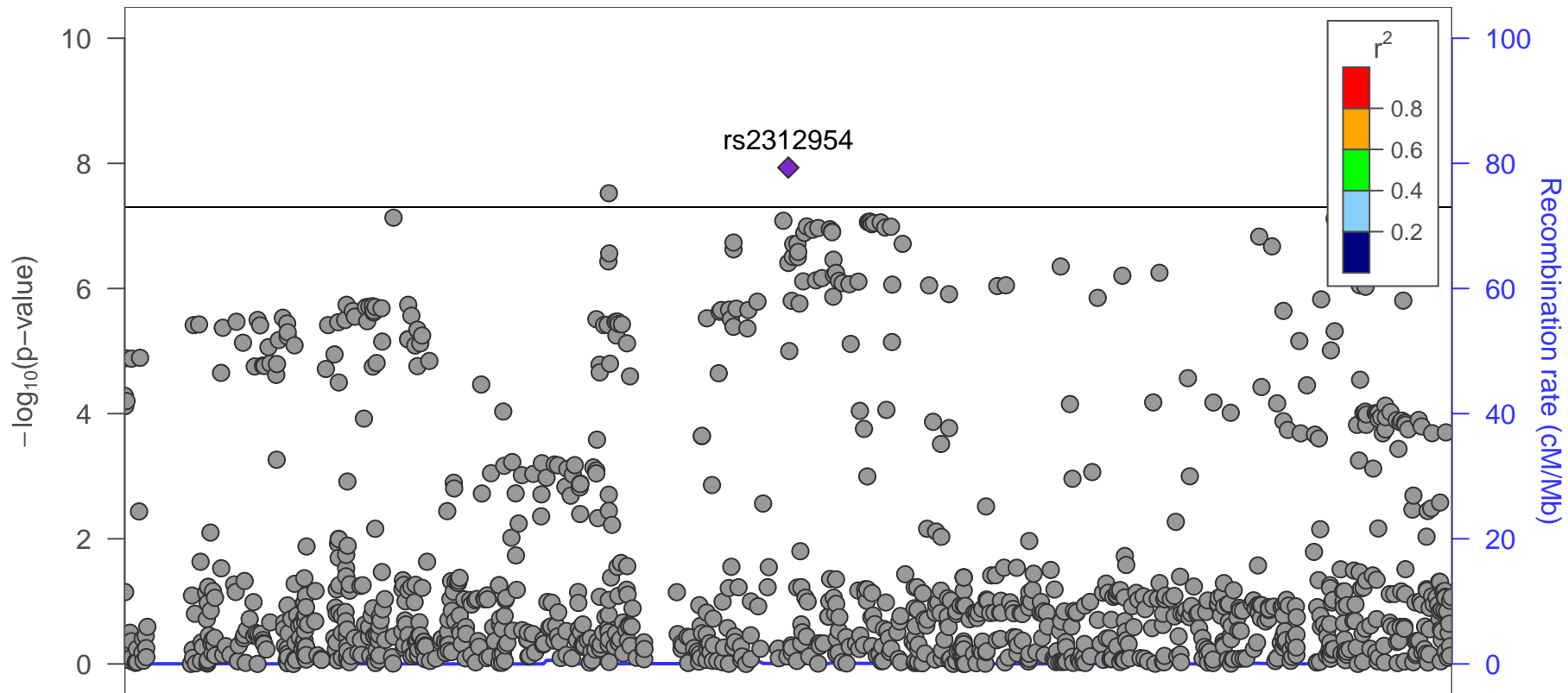




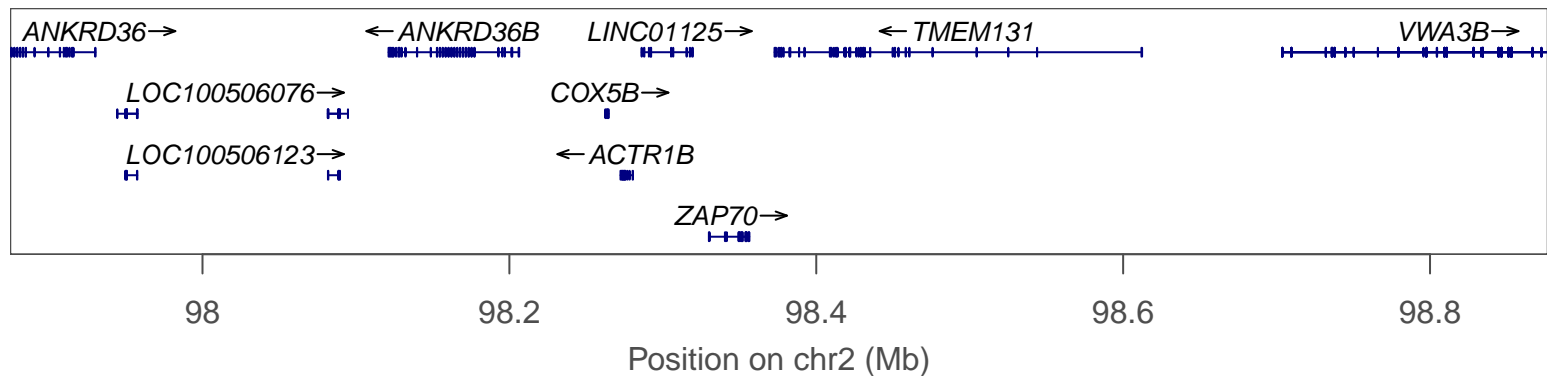
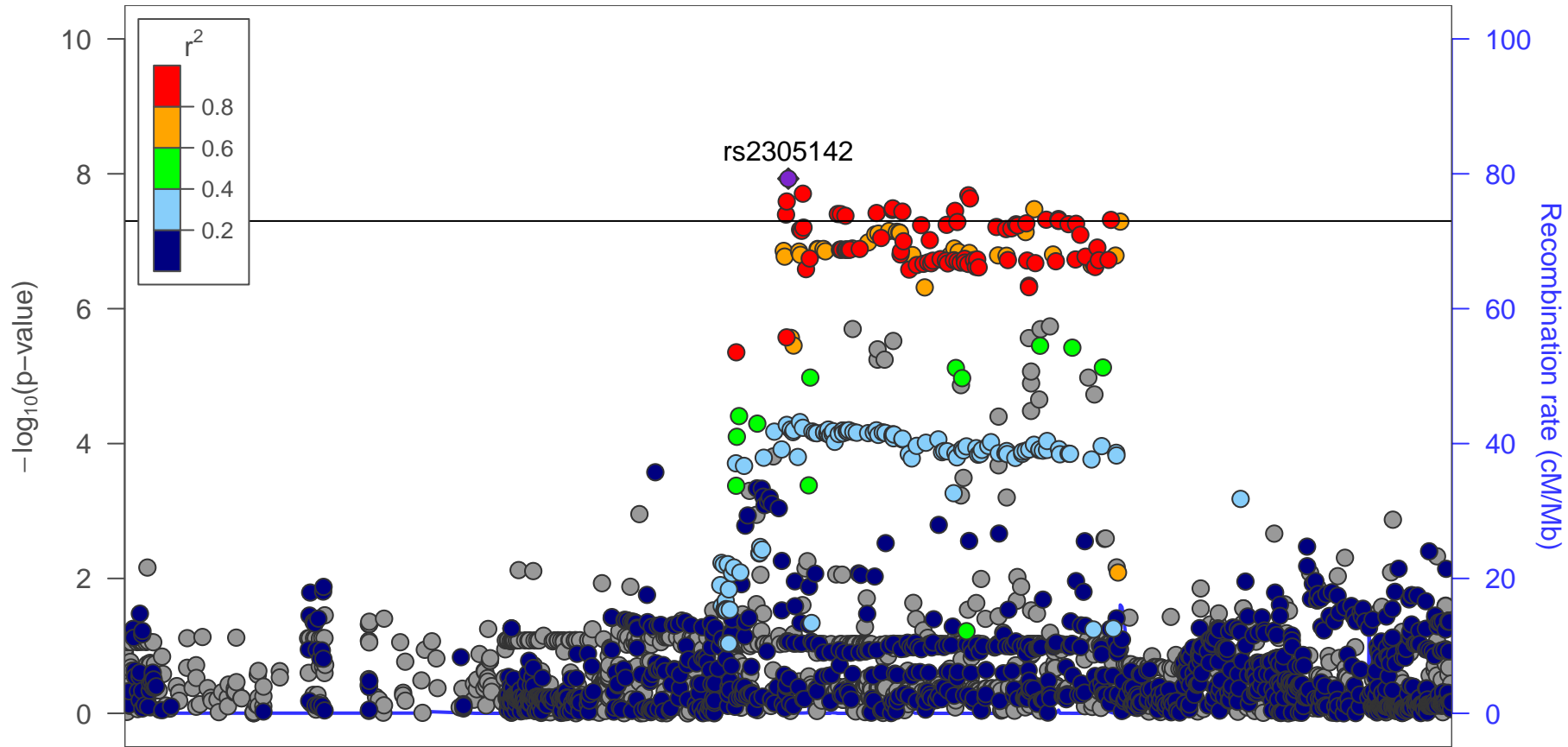


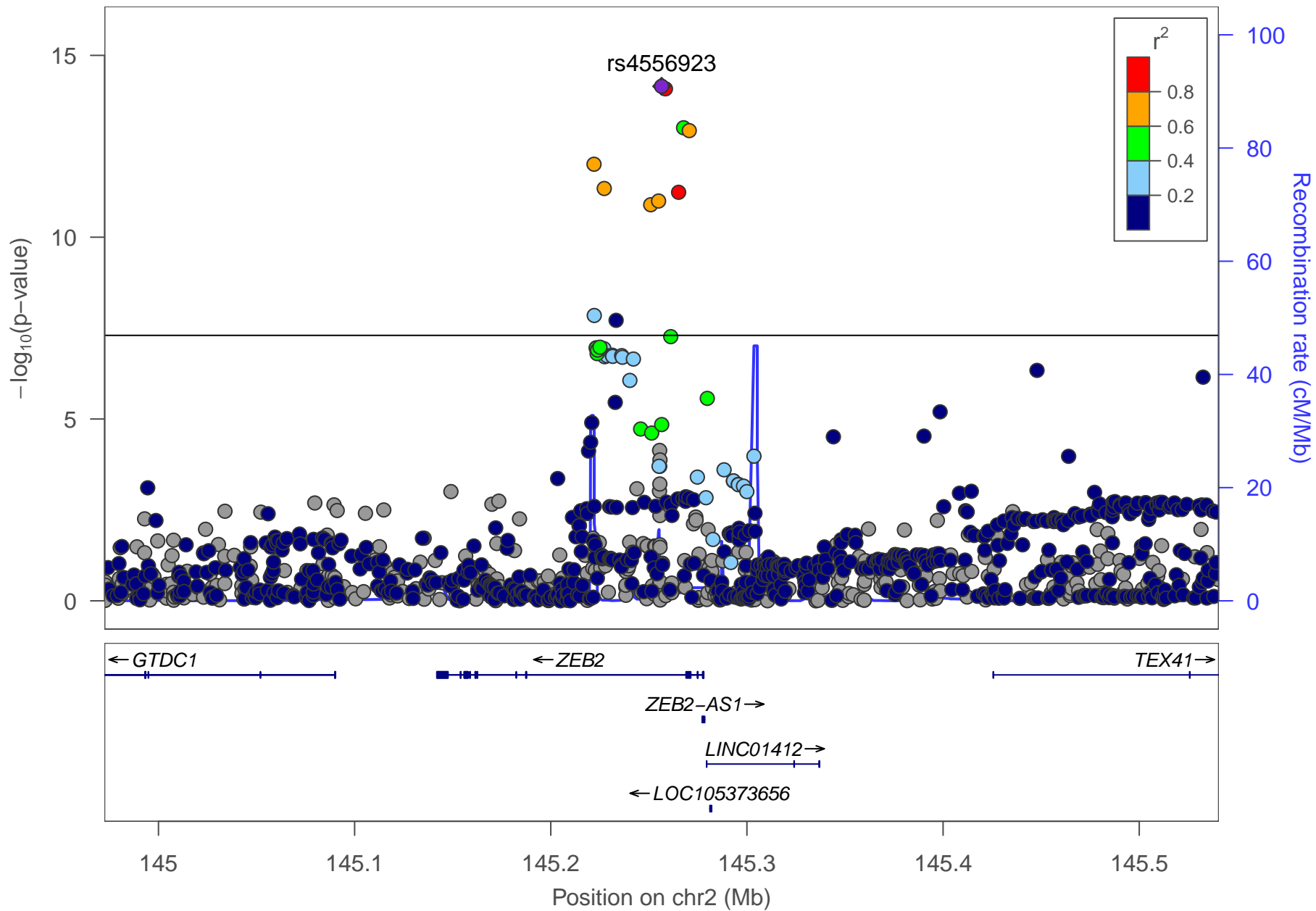


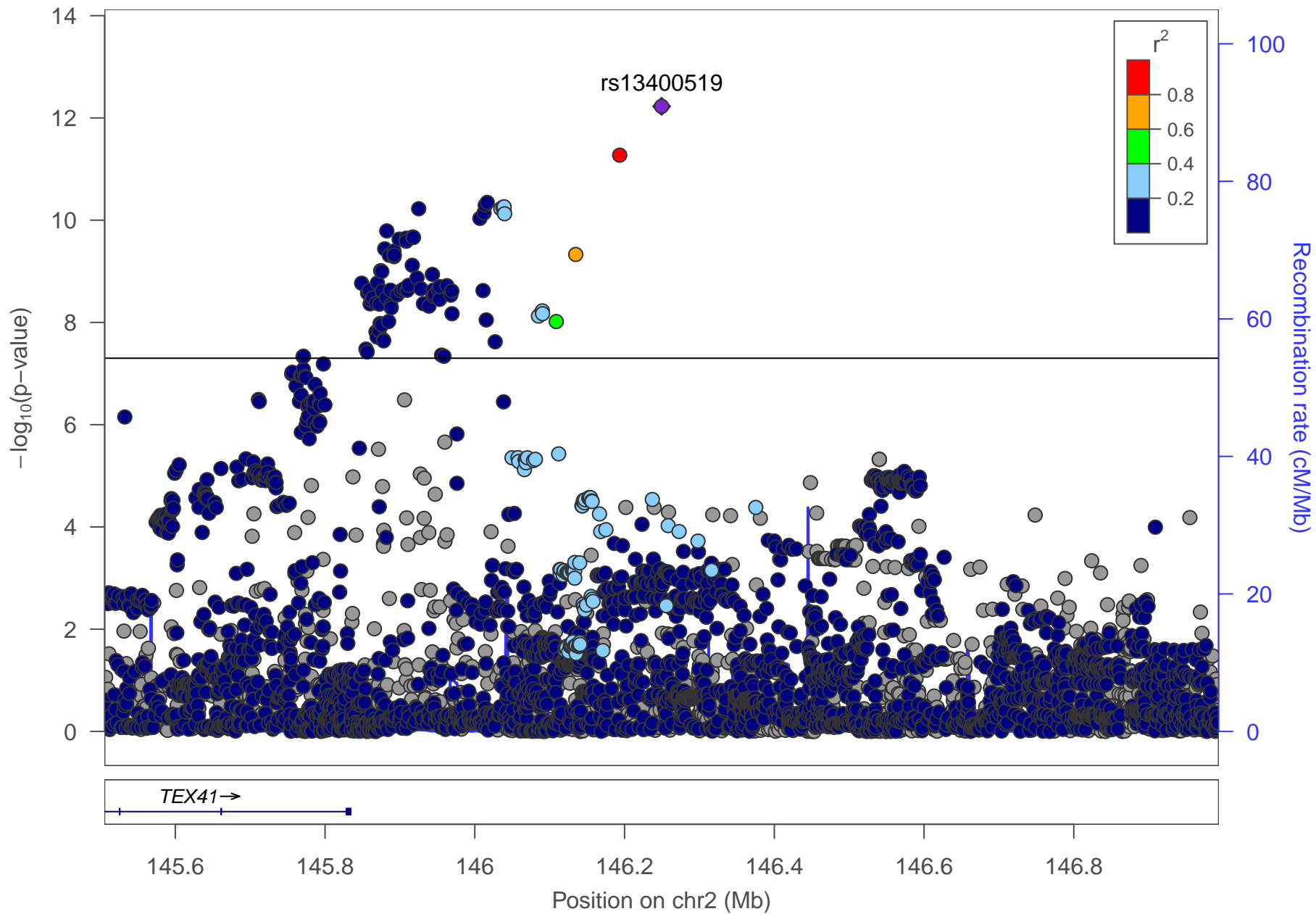


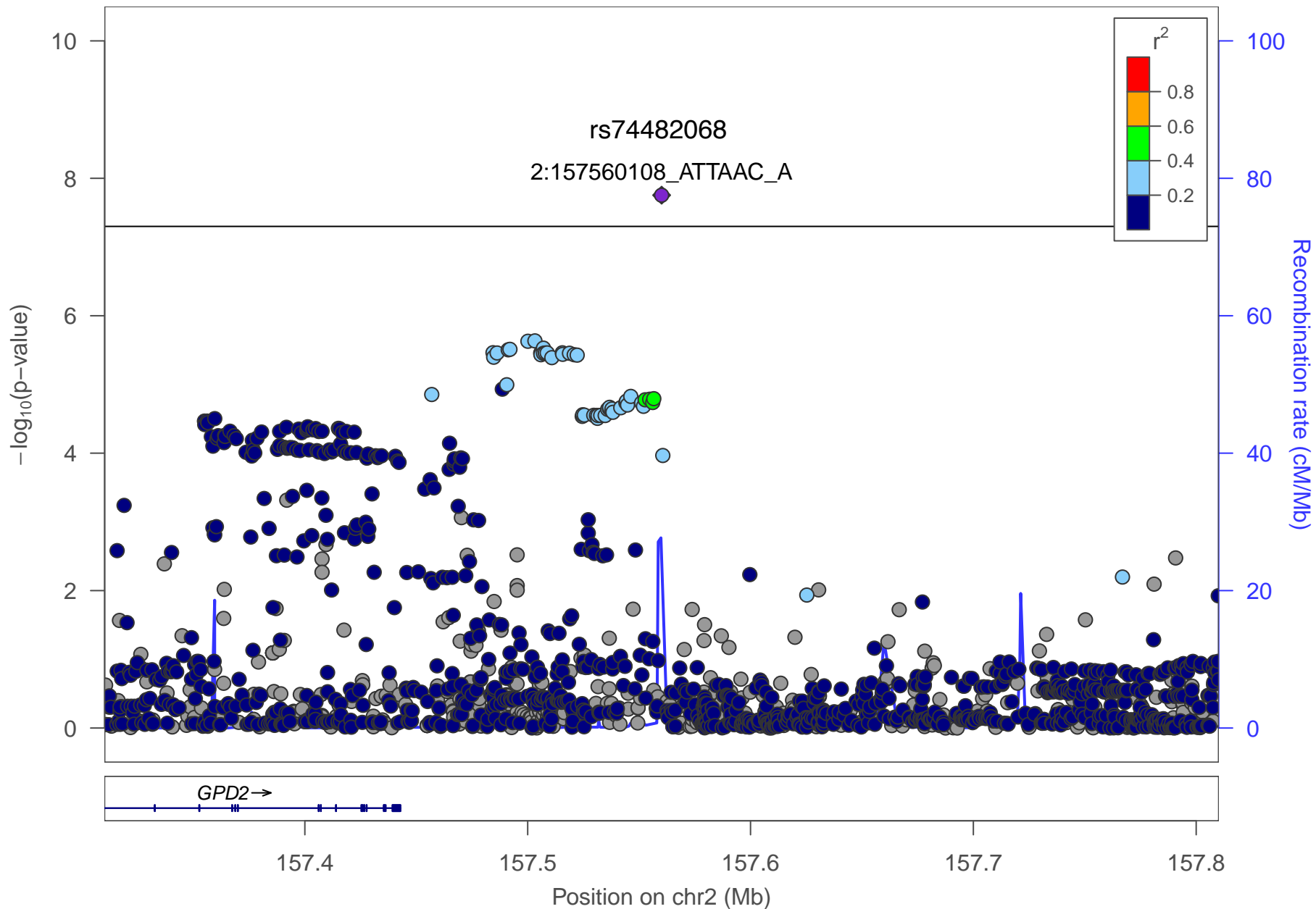


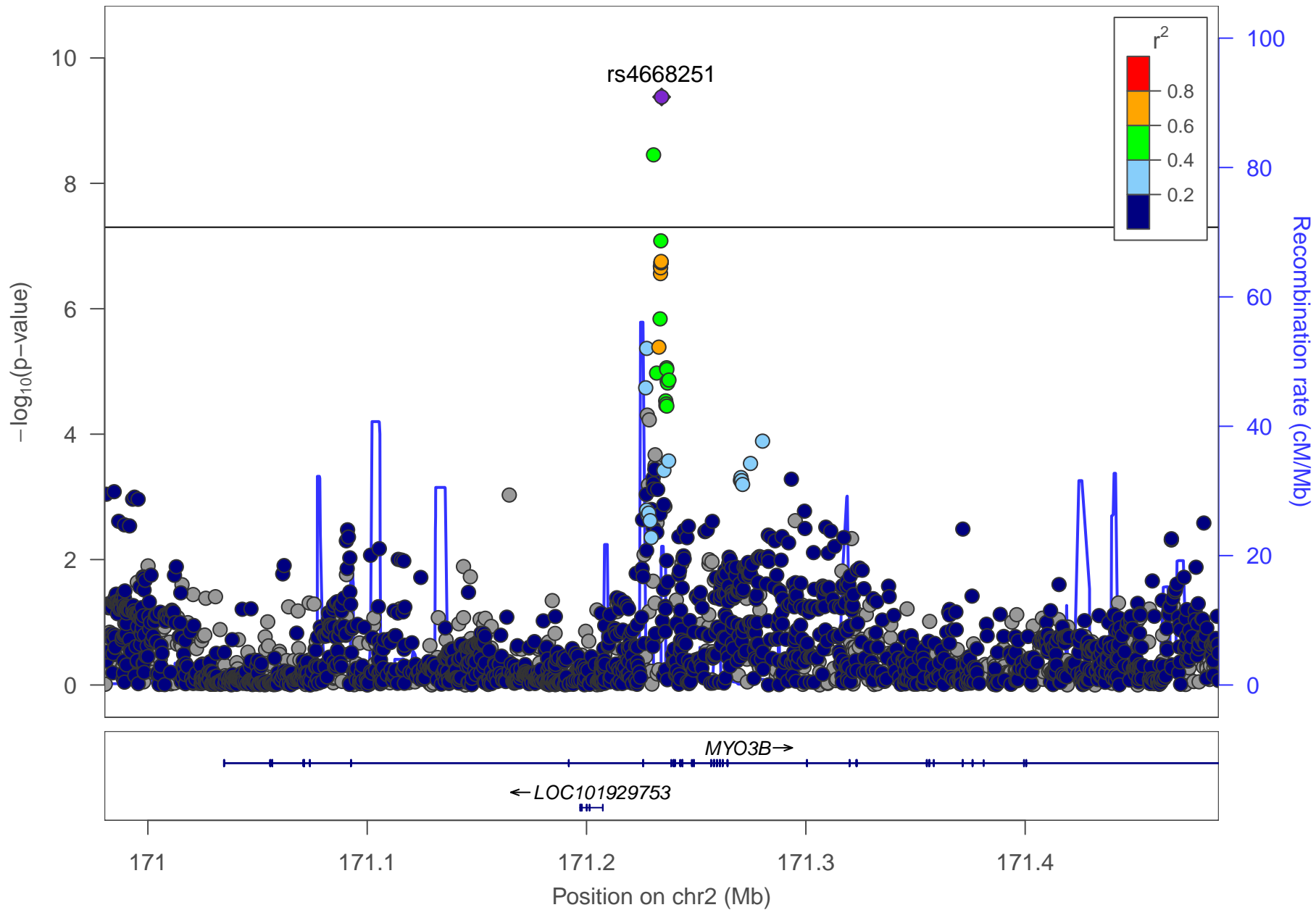


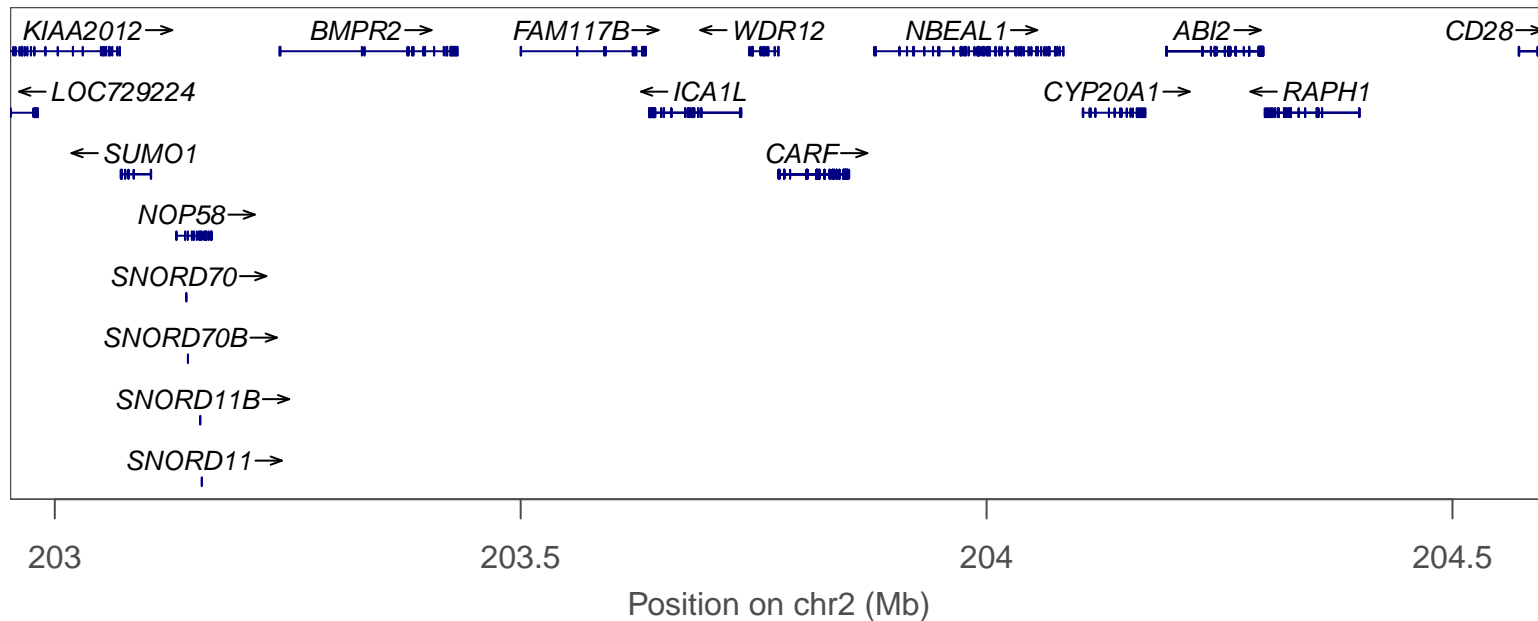
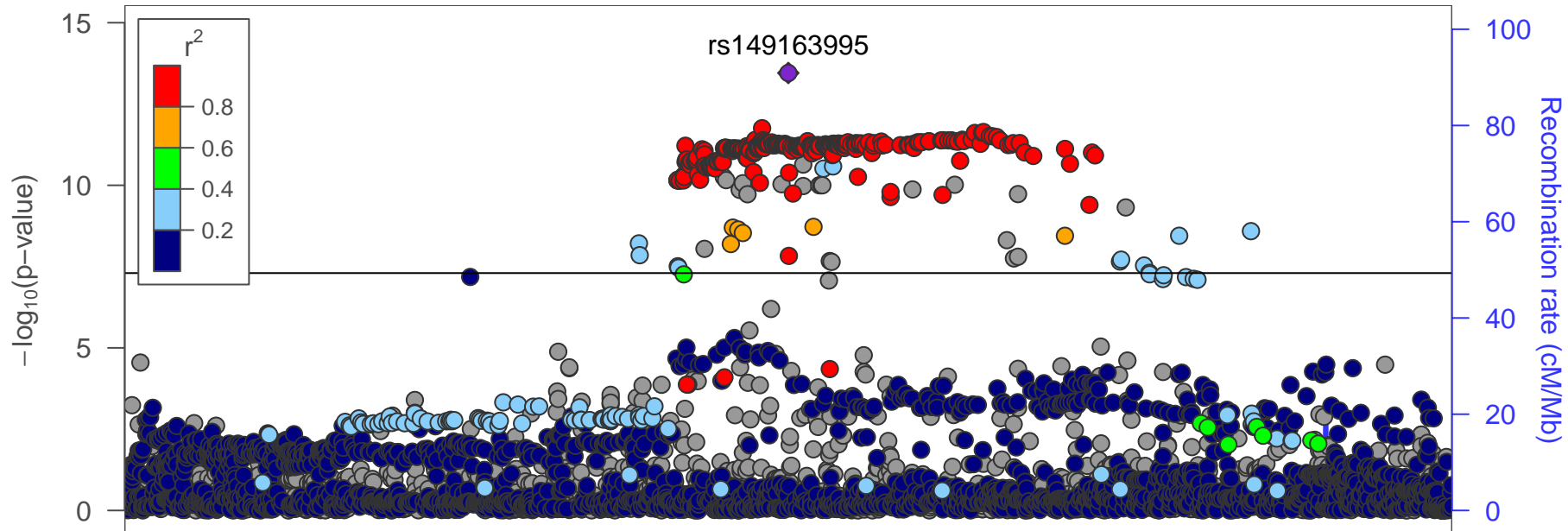


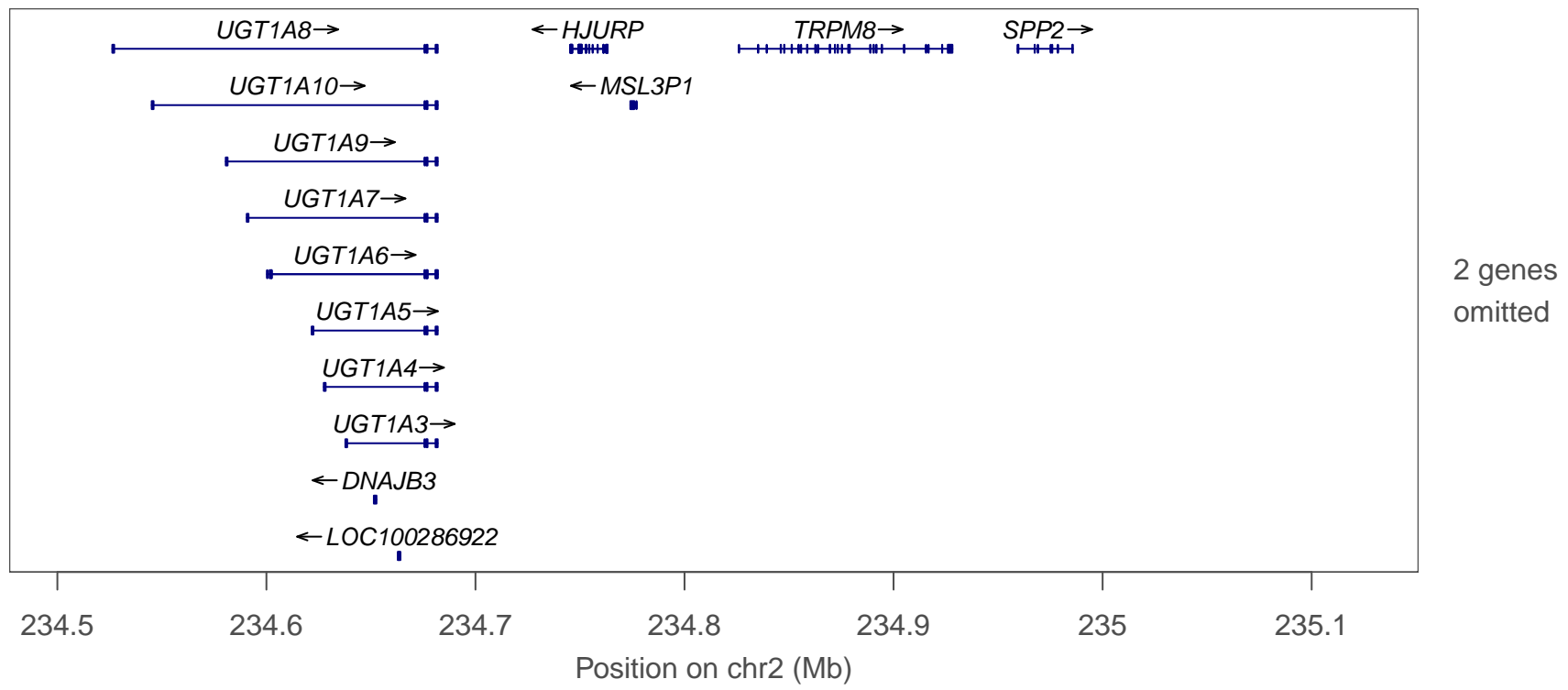
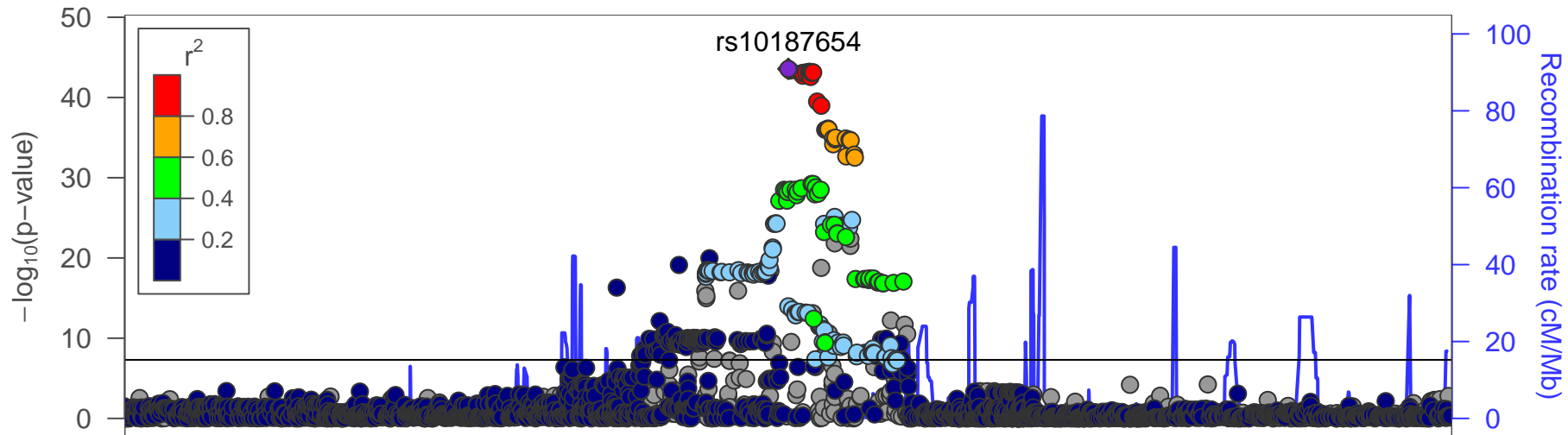


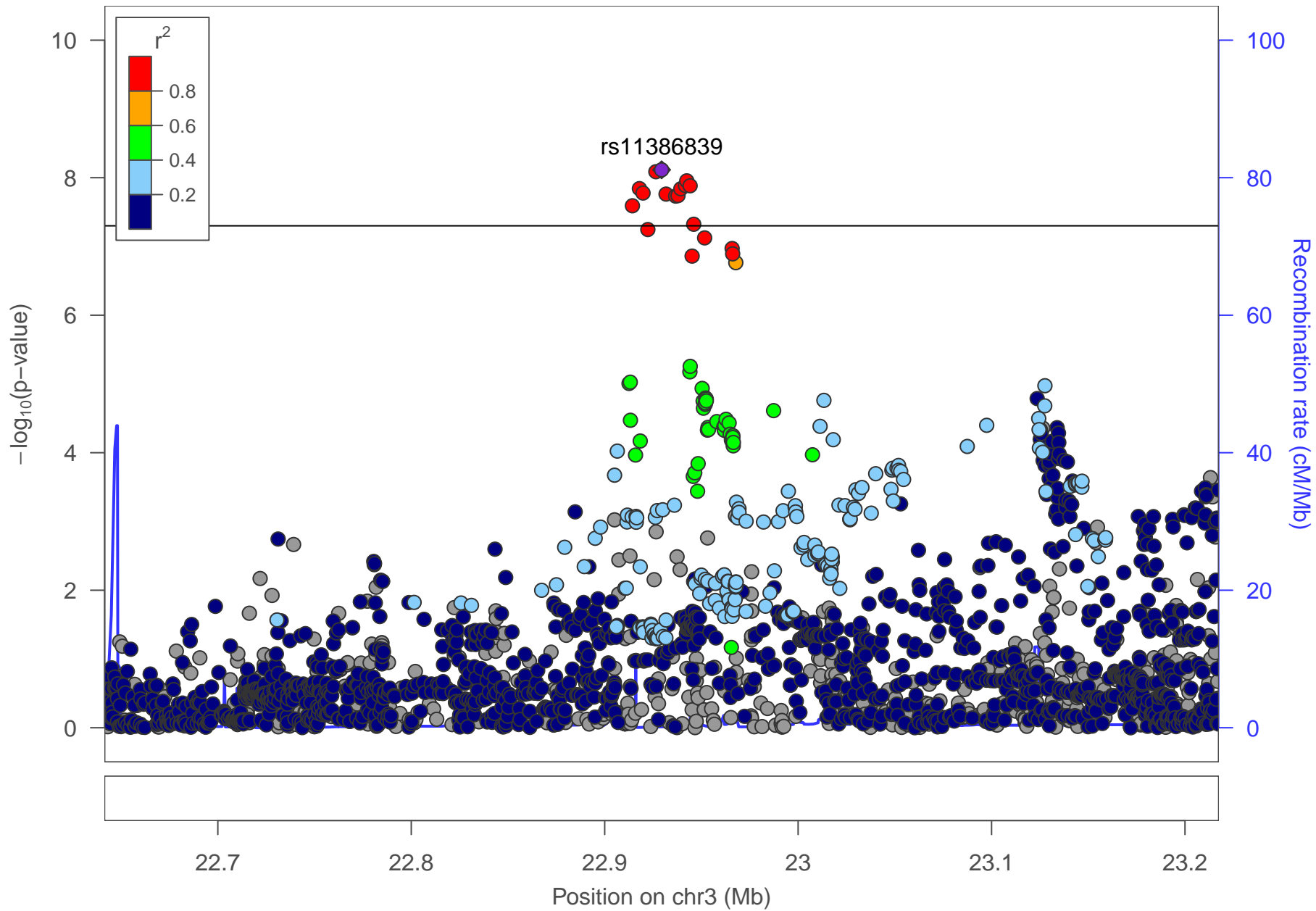




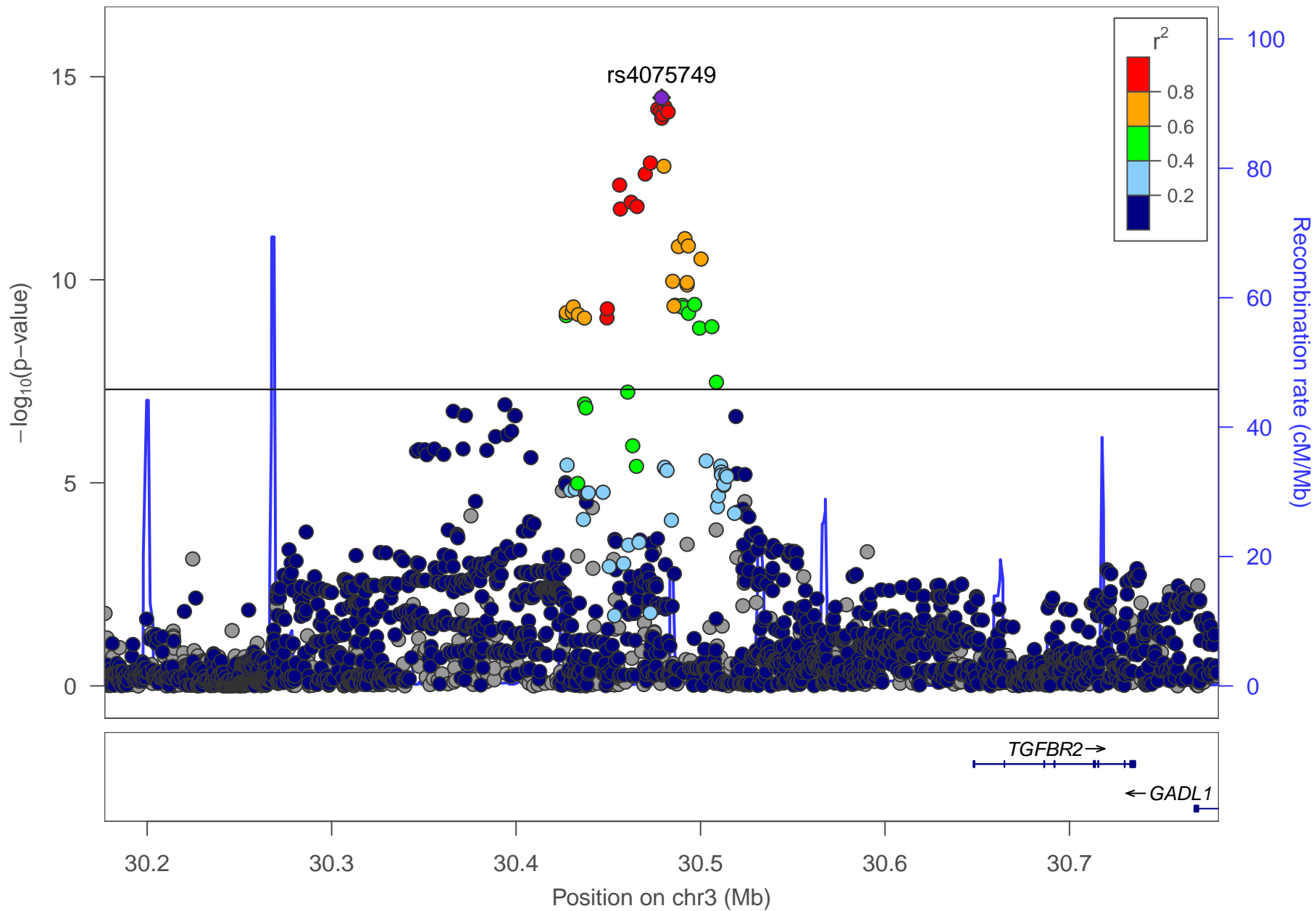


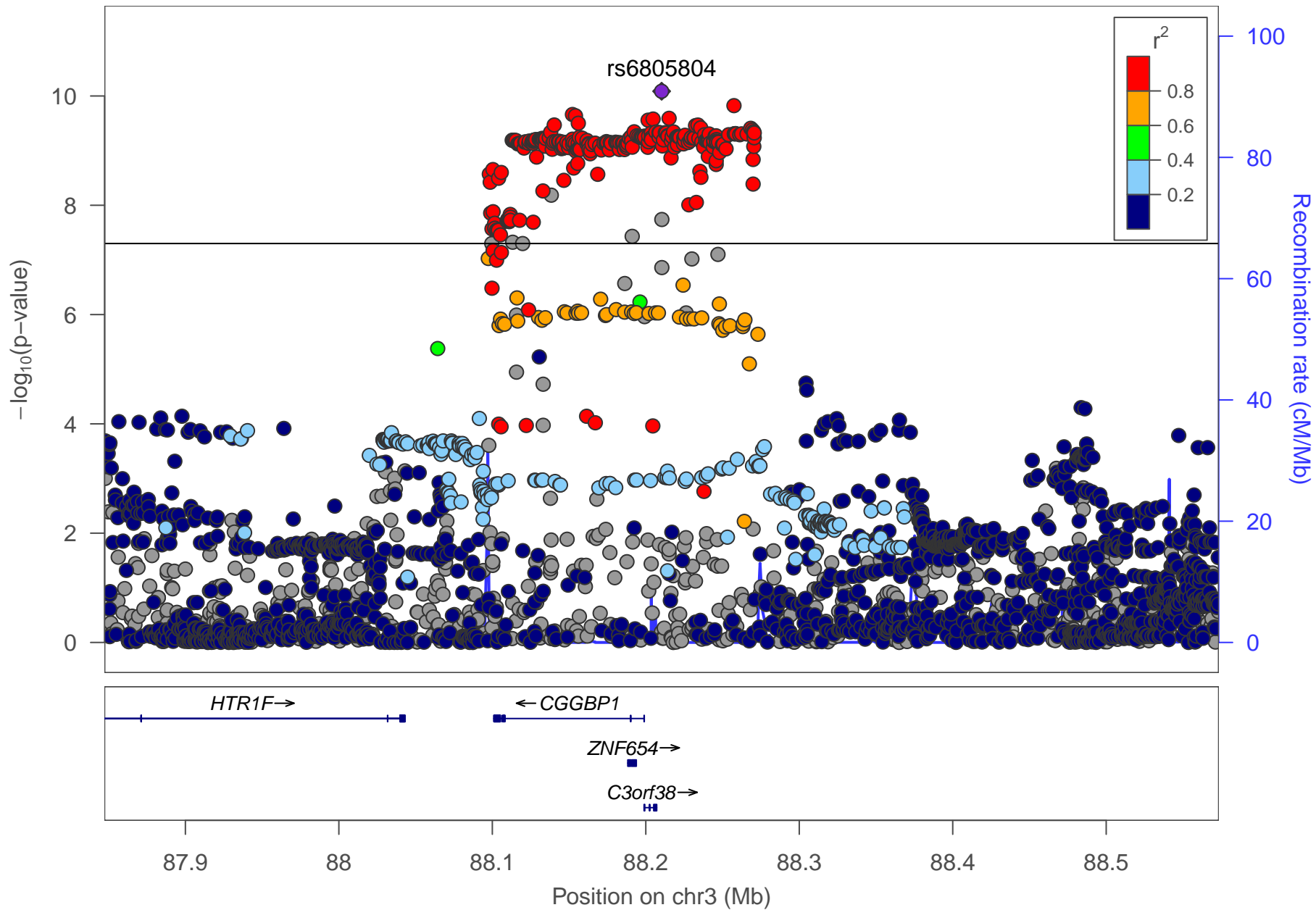


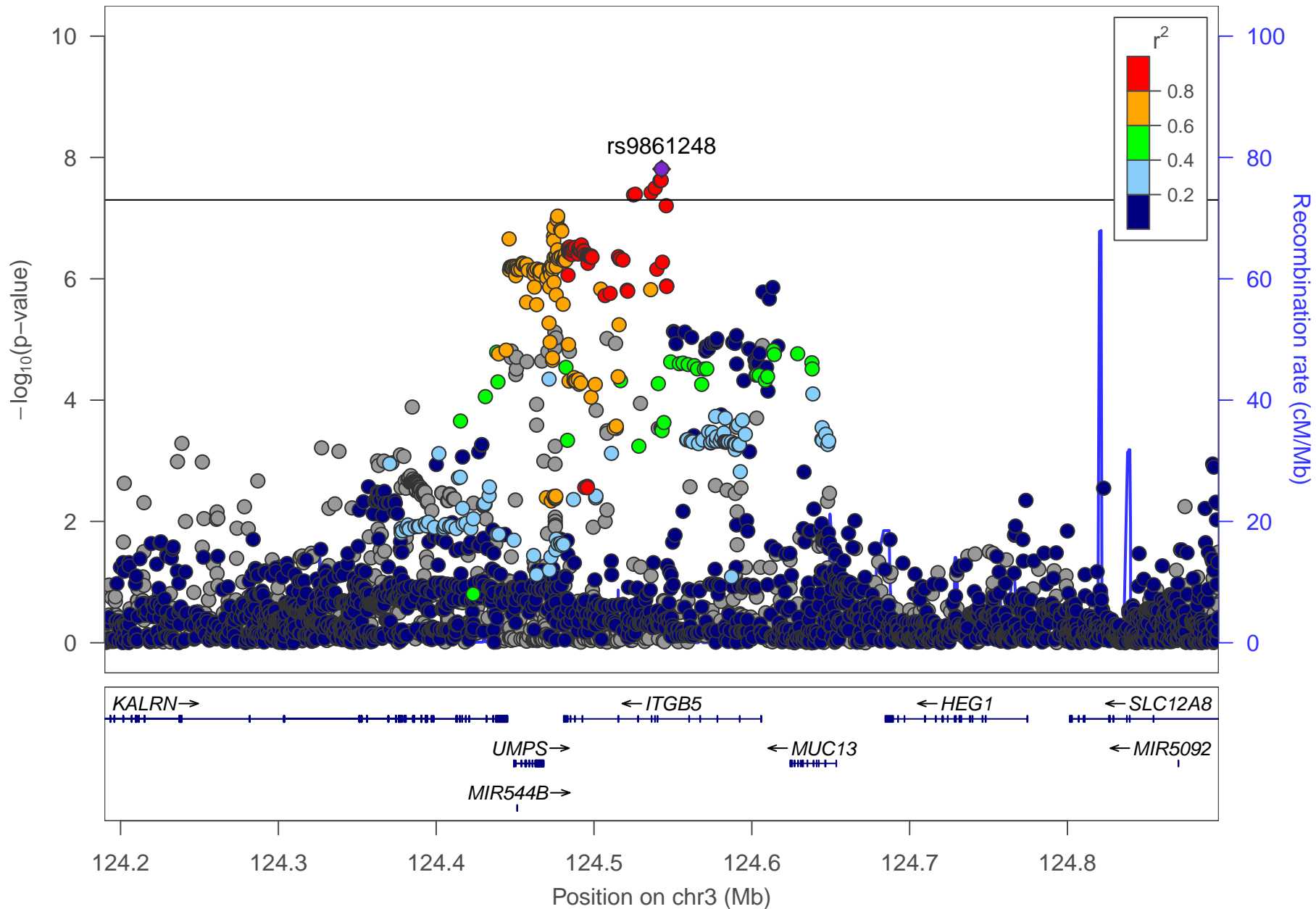


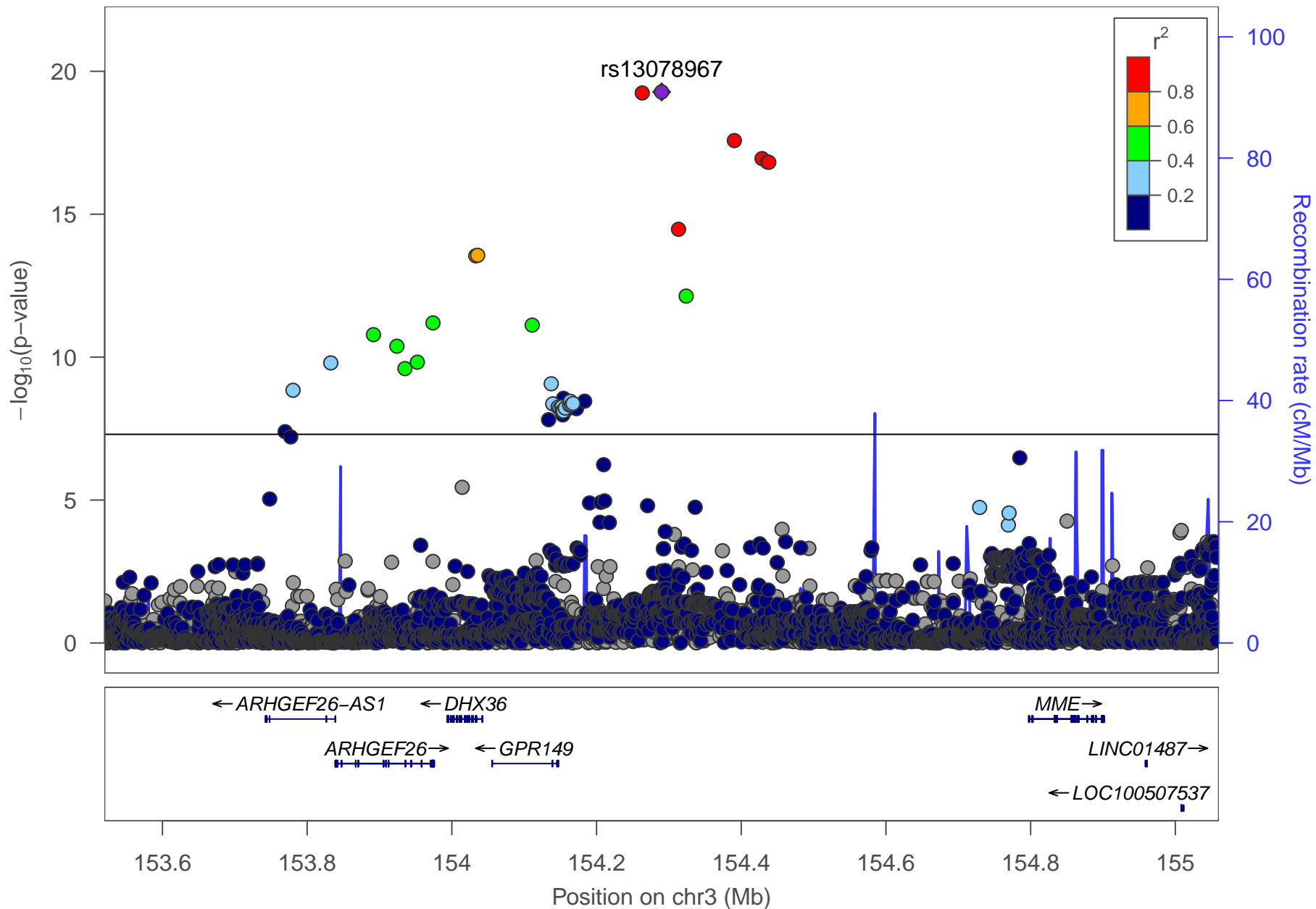


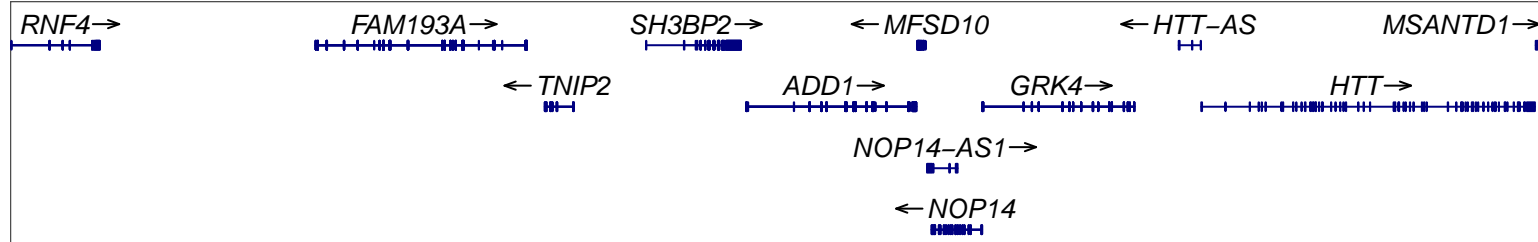
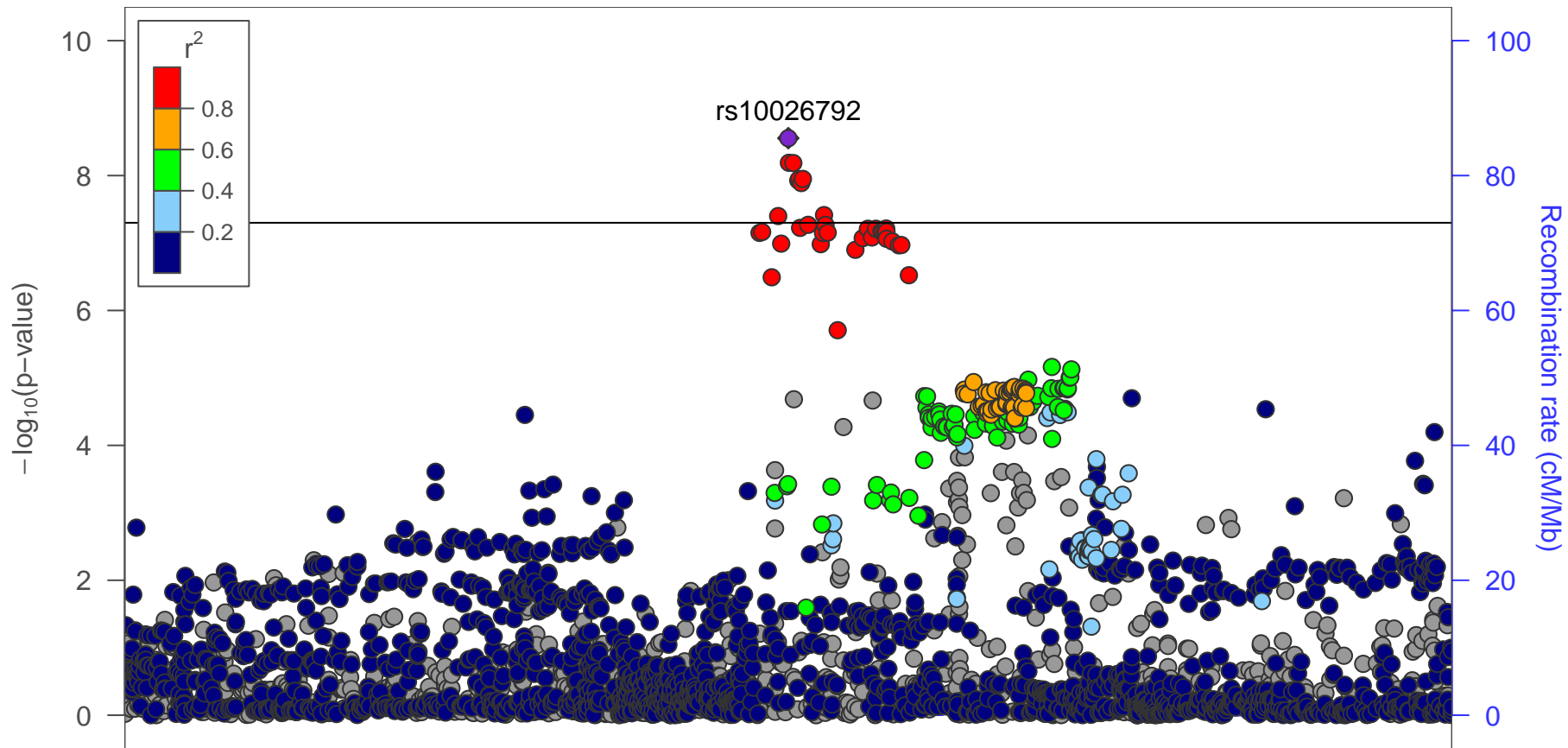












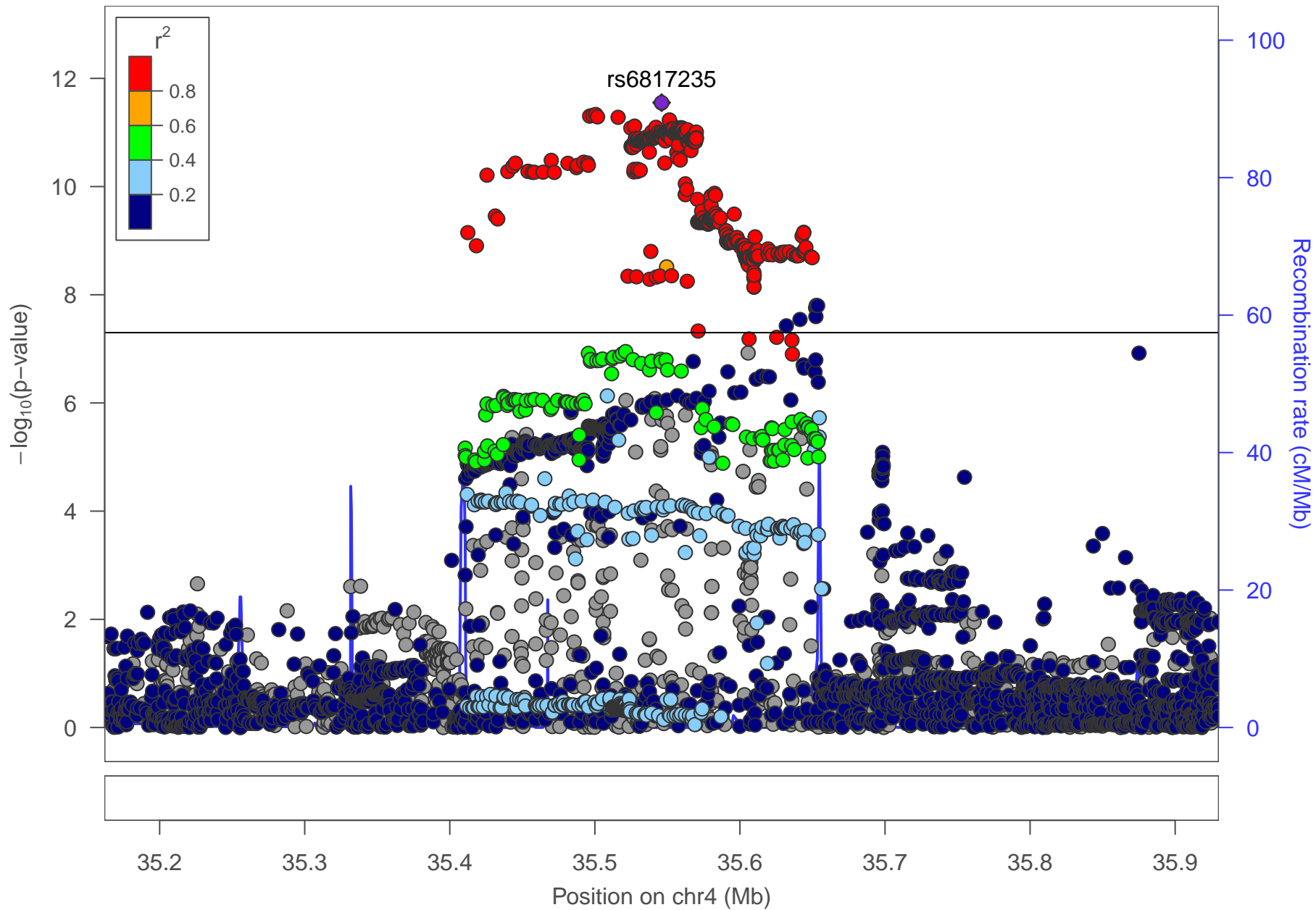
2.6

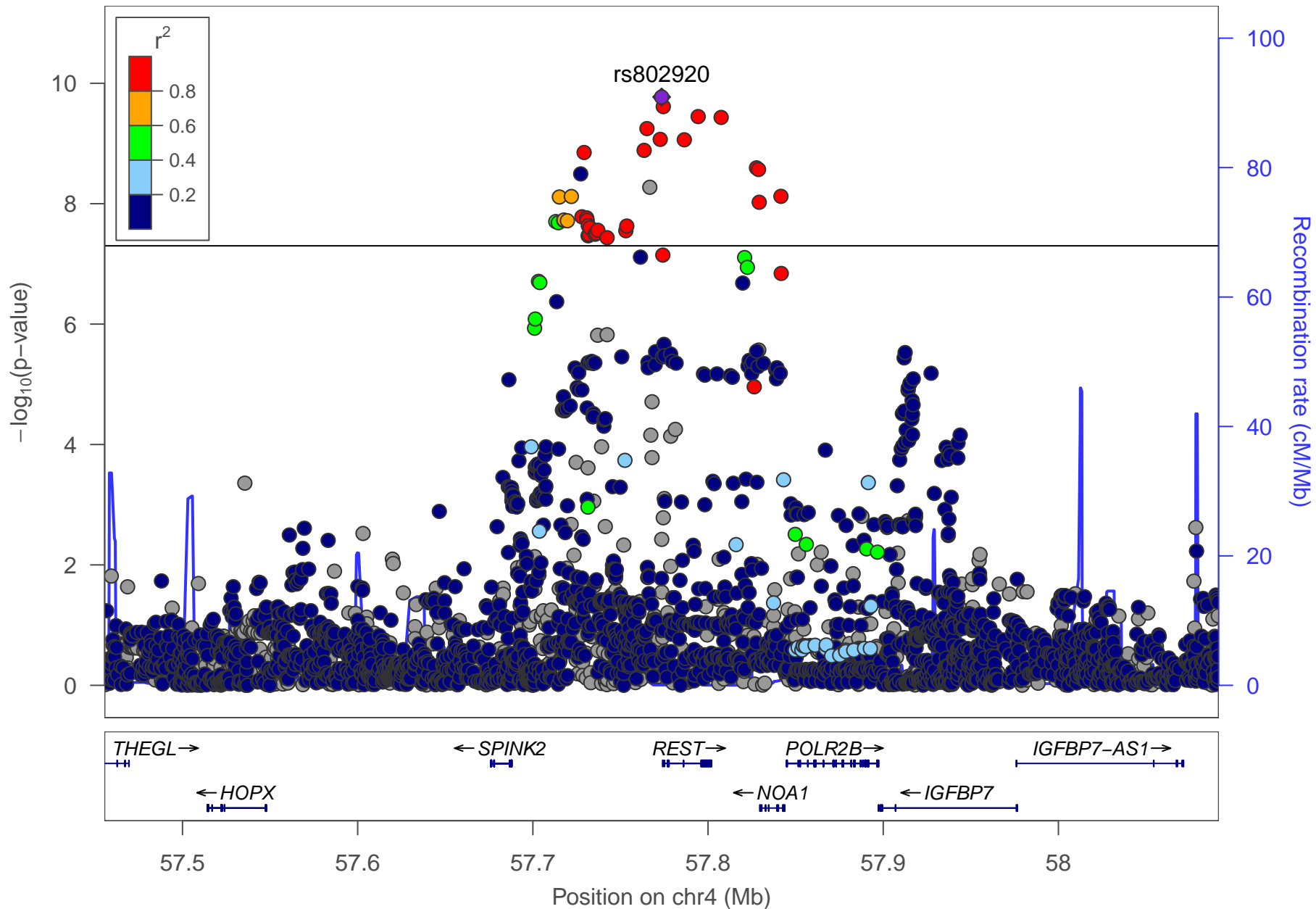
2.8

3

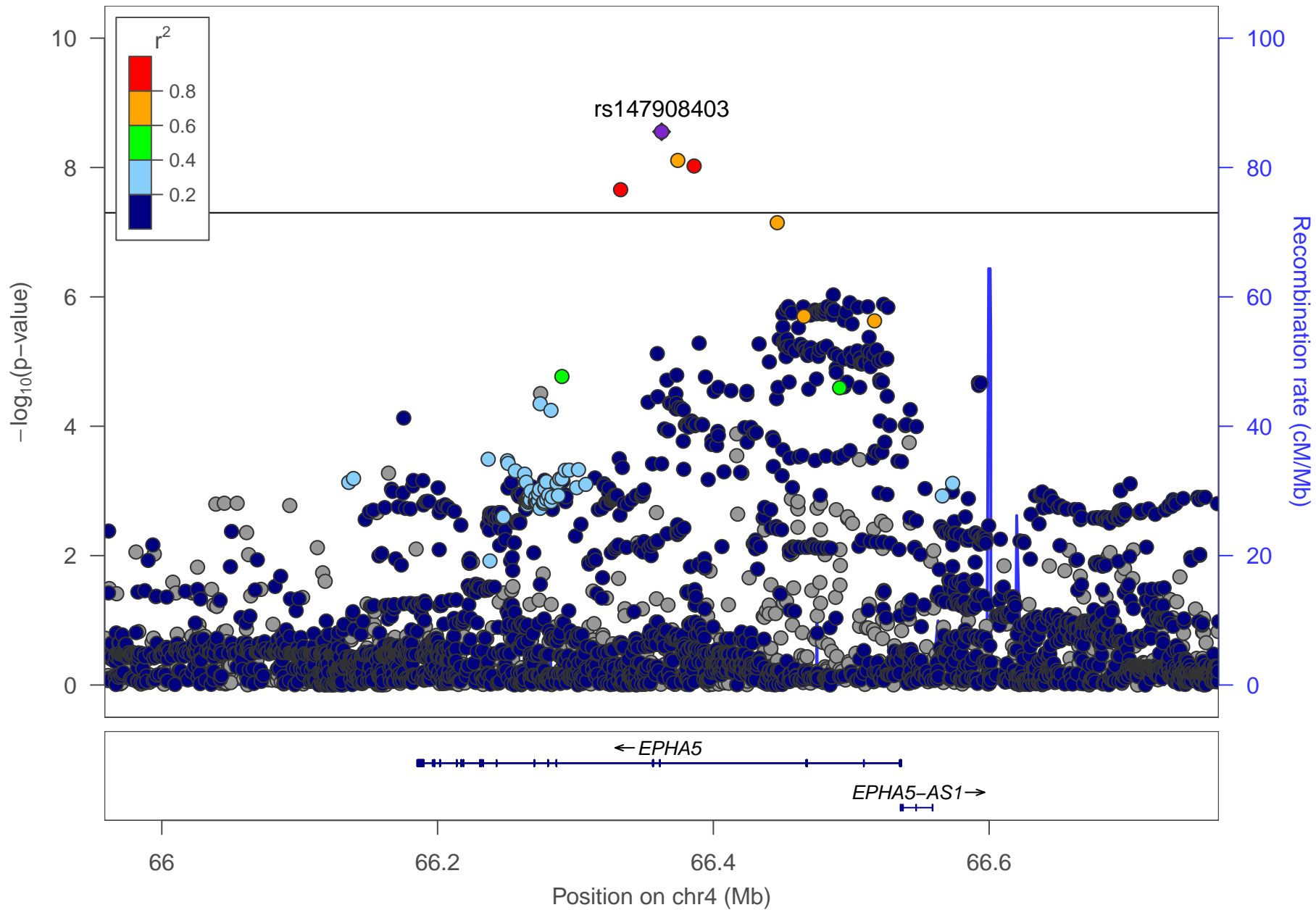
3.2

Position on chr4 (Mb)

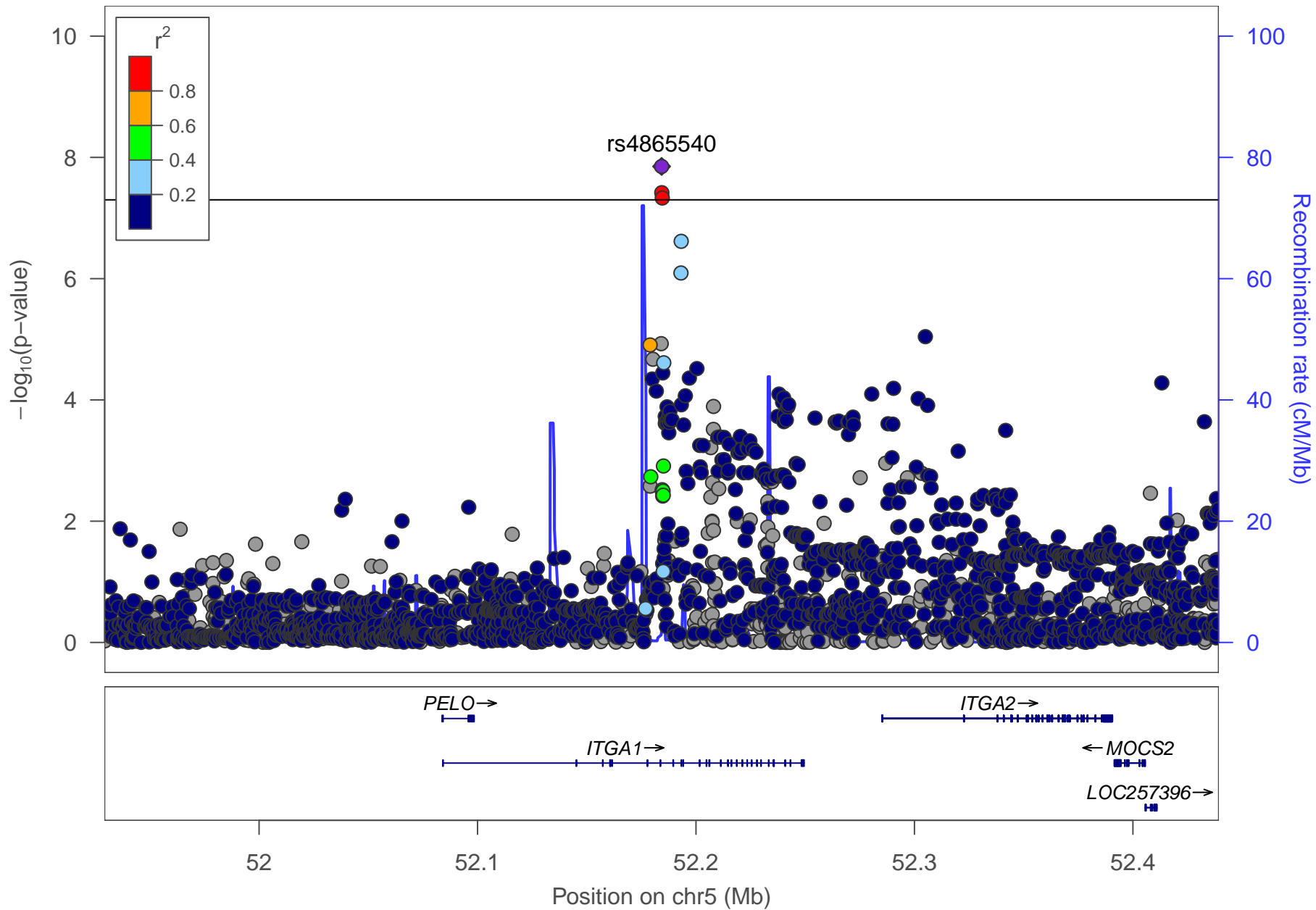


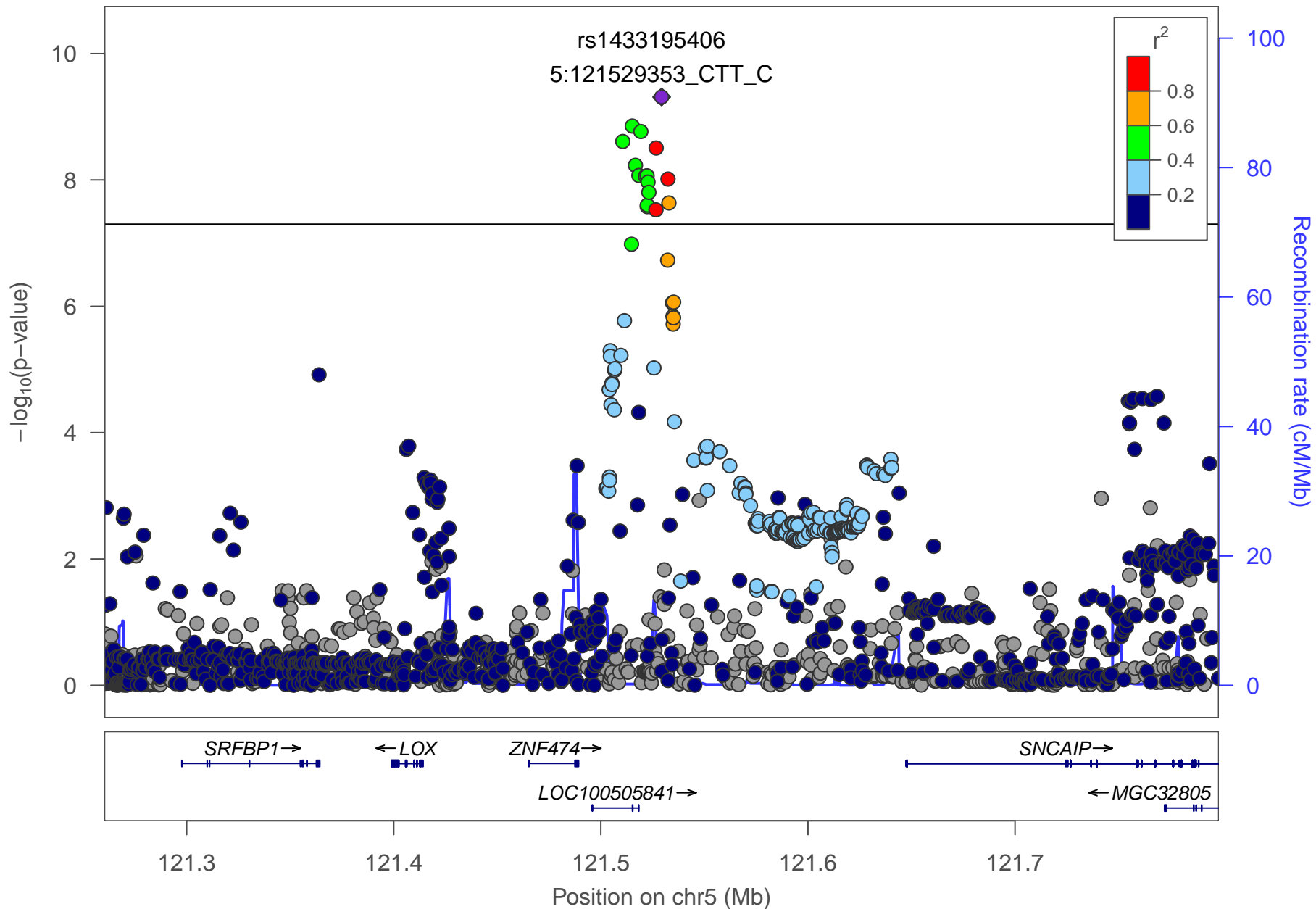


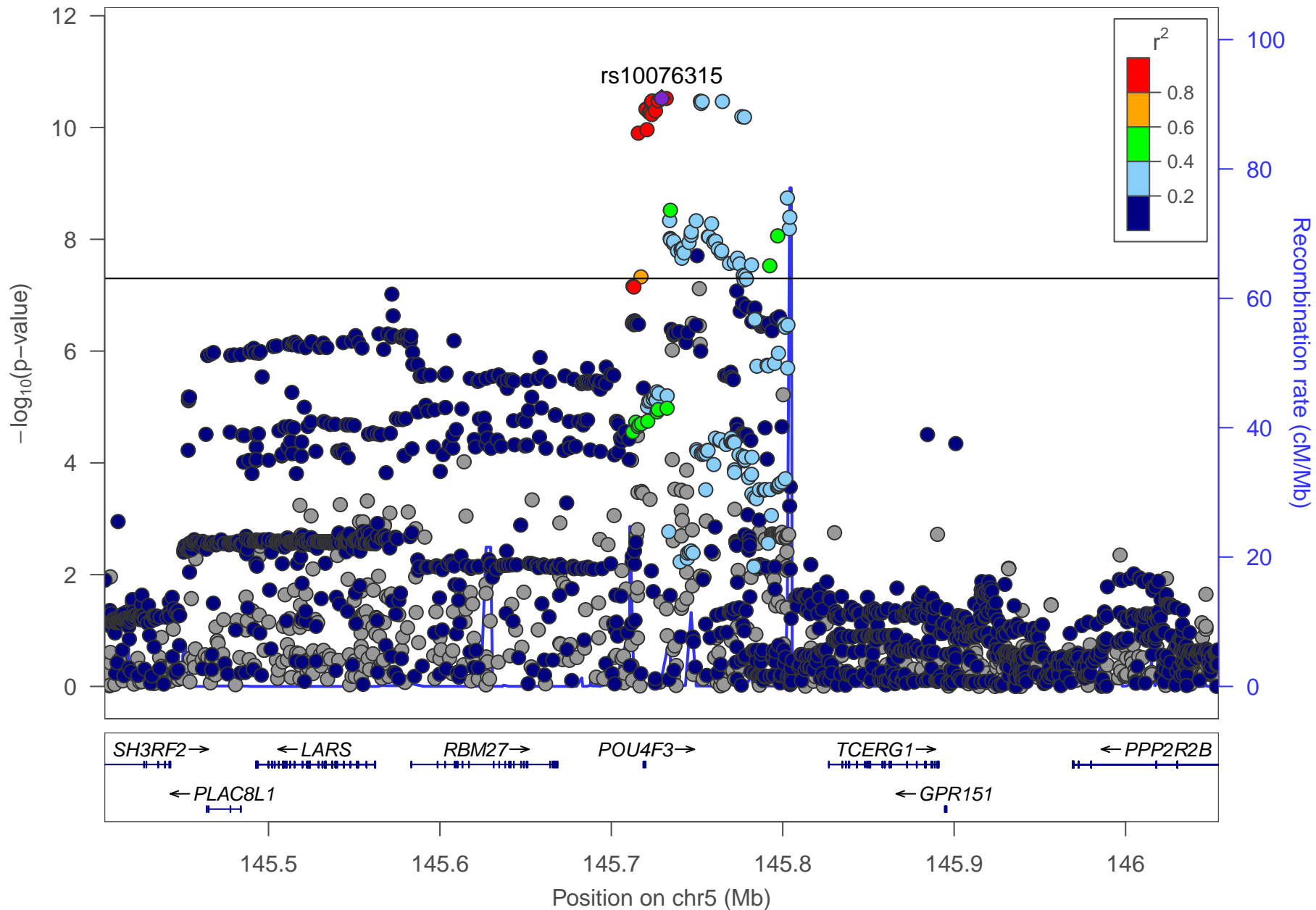


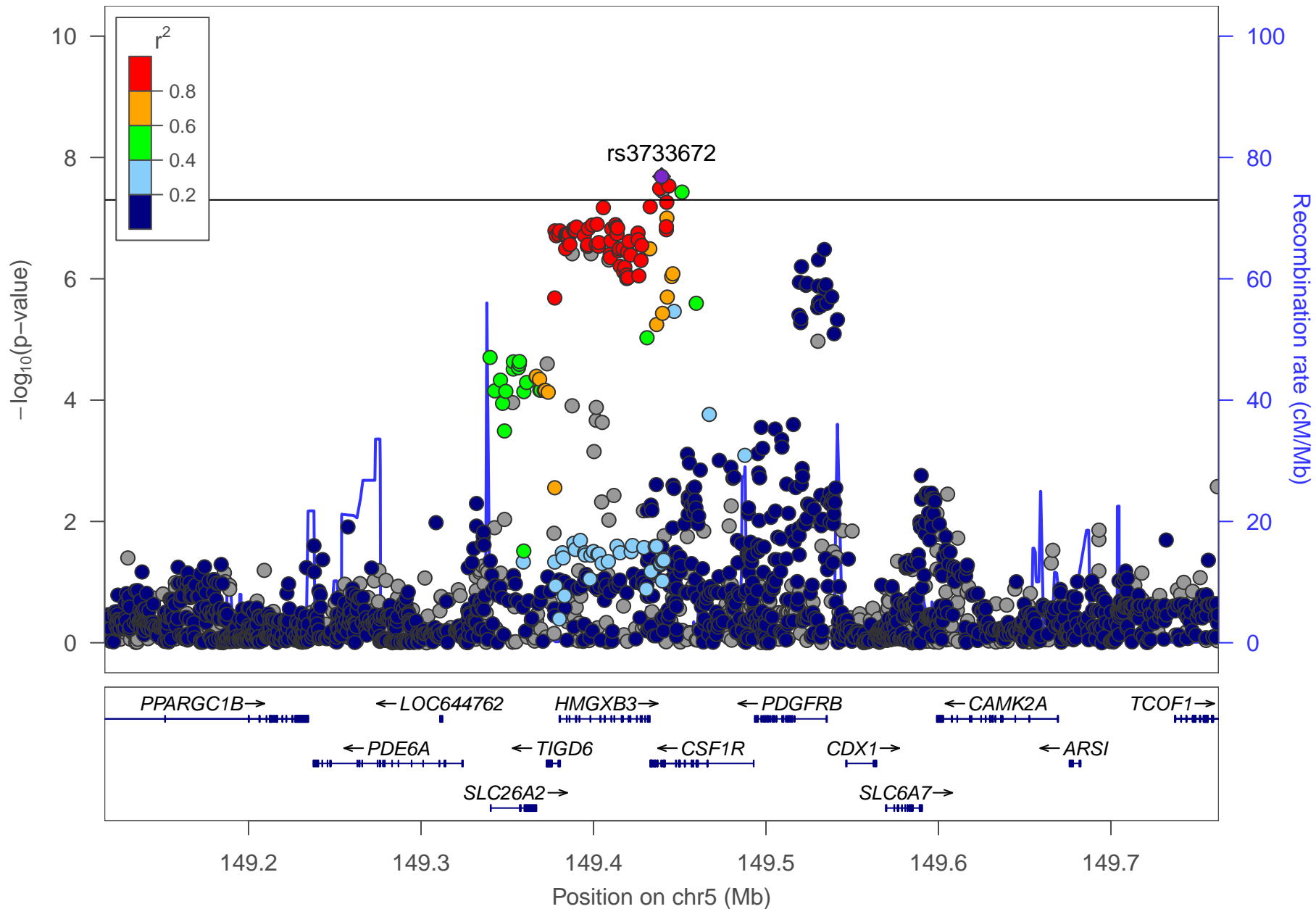


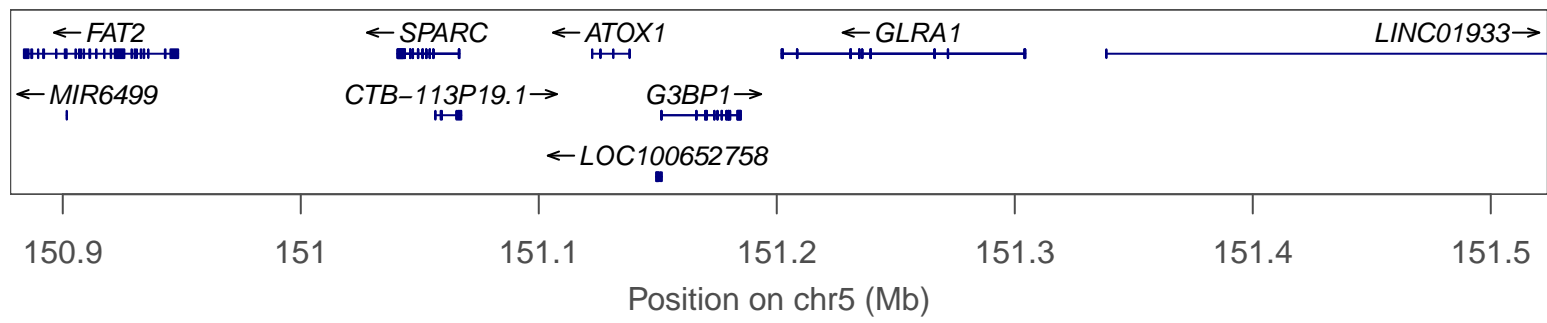
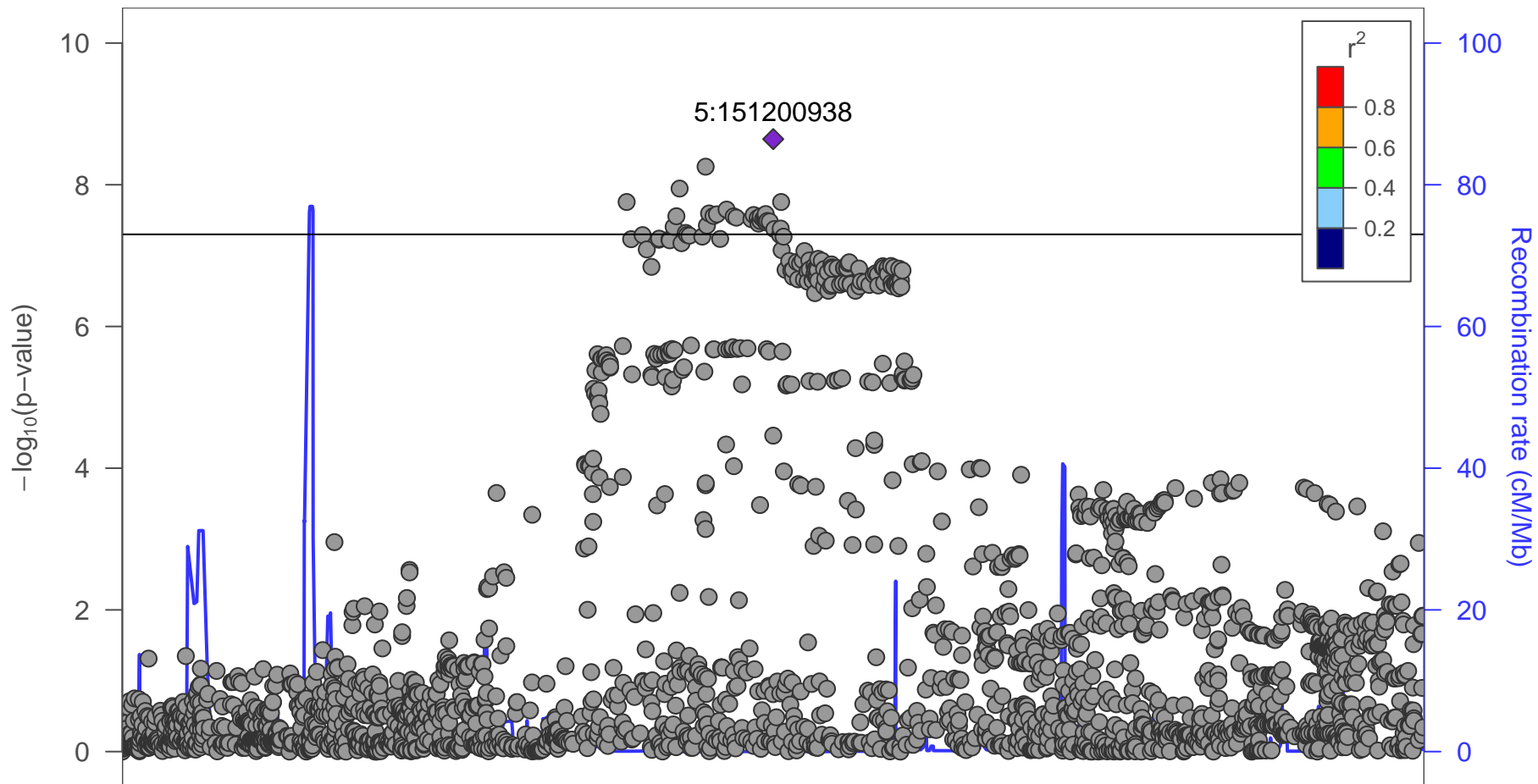


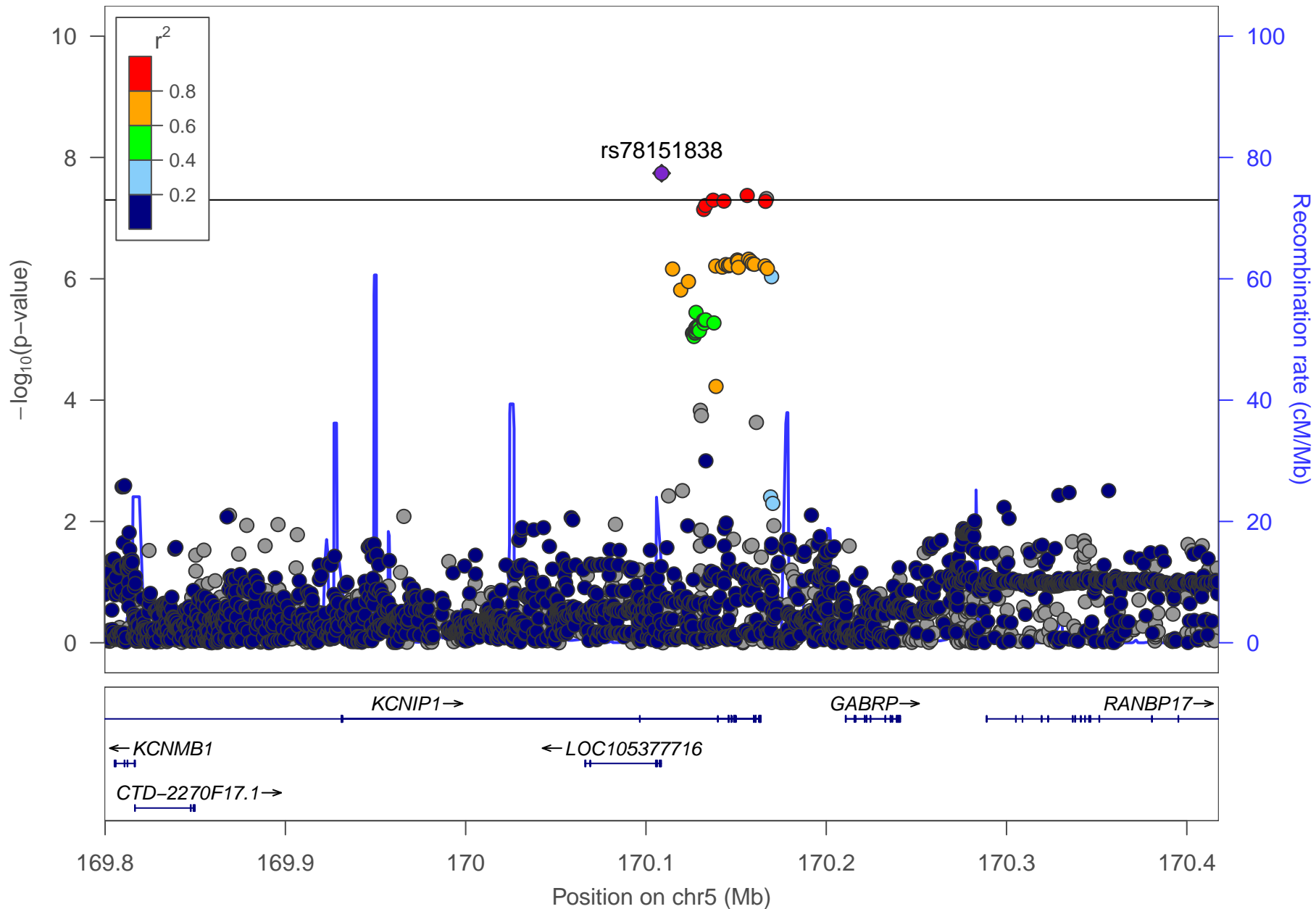


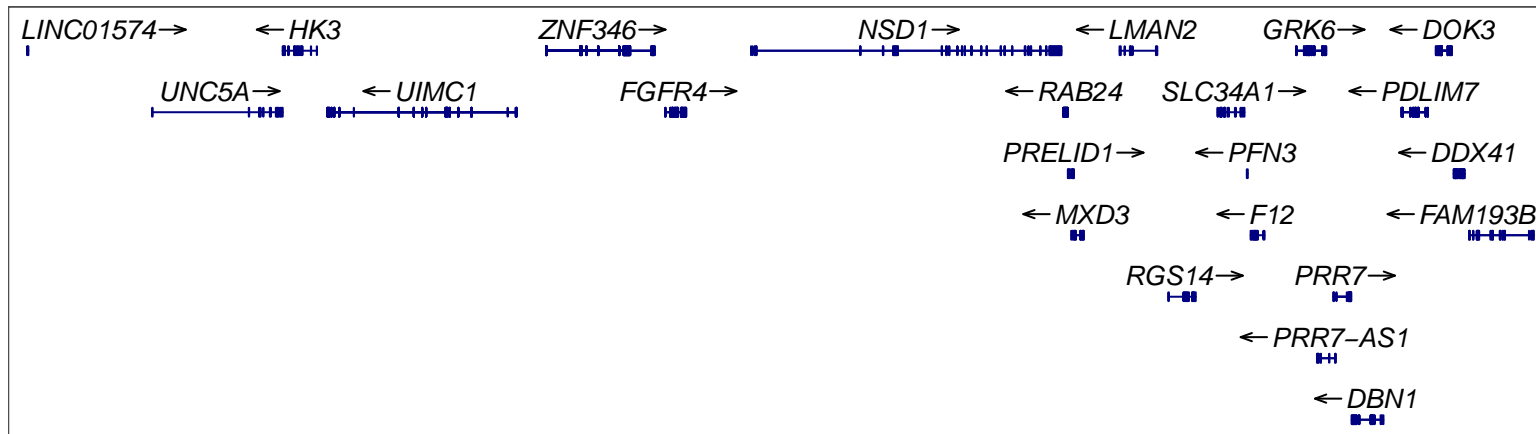
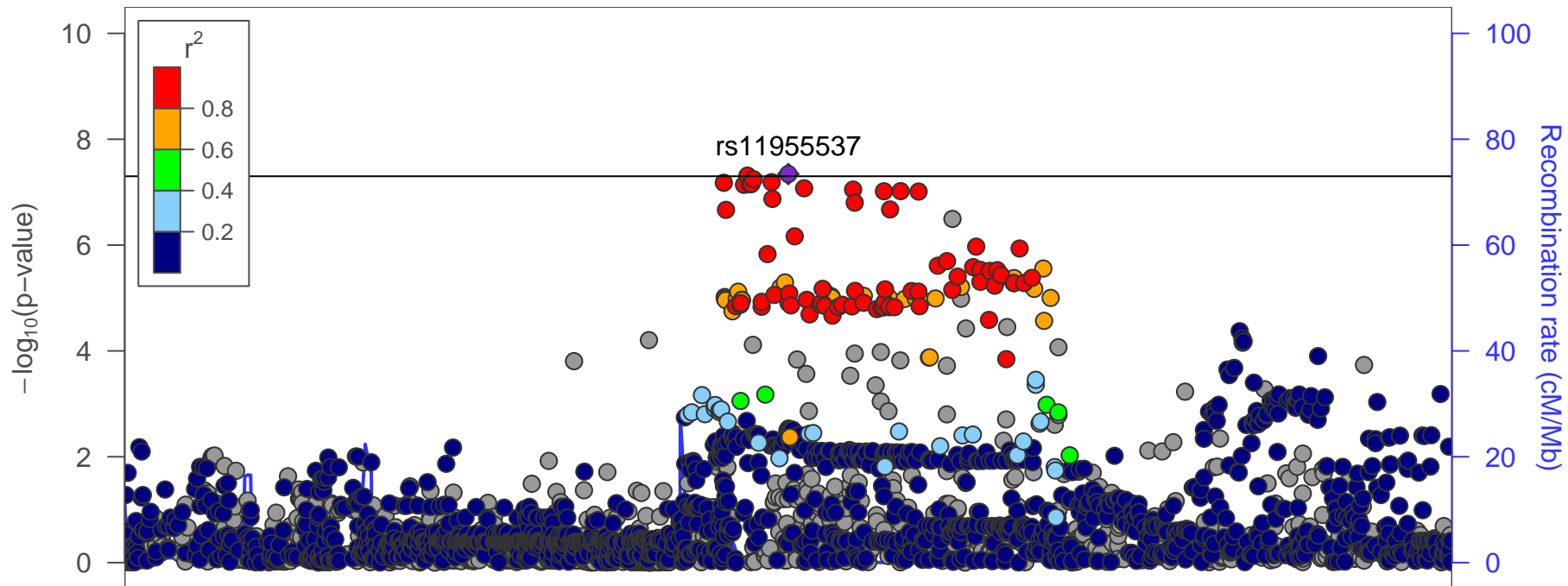












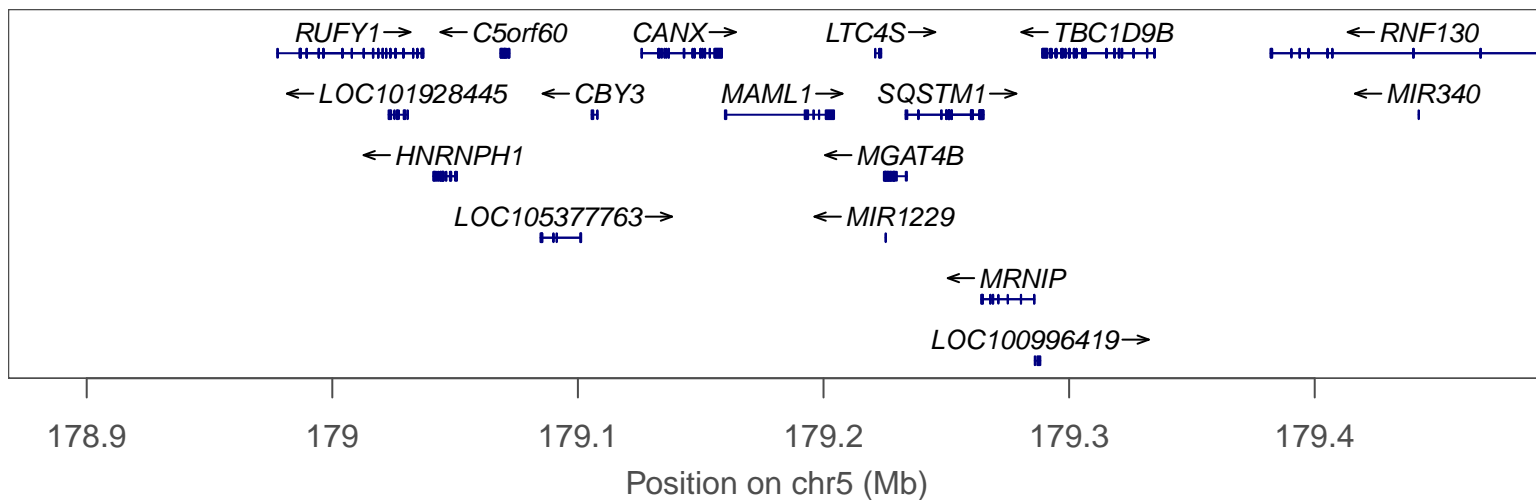
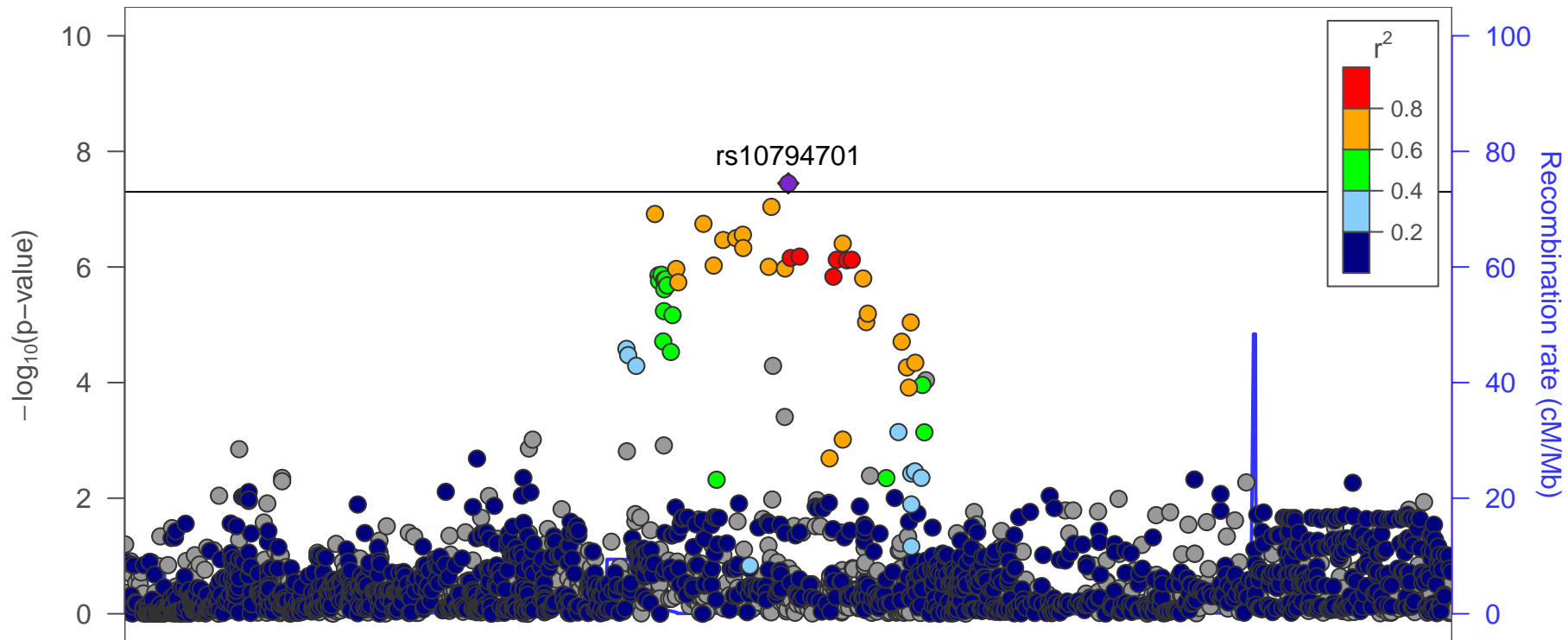
176.2

176.4

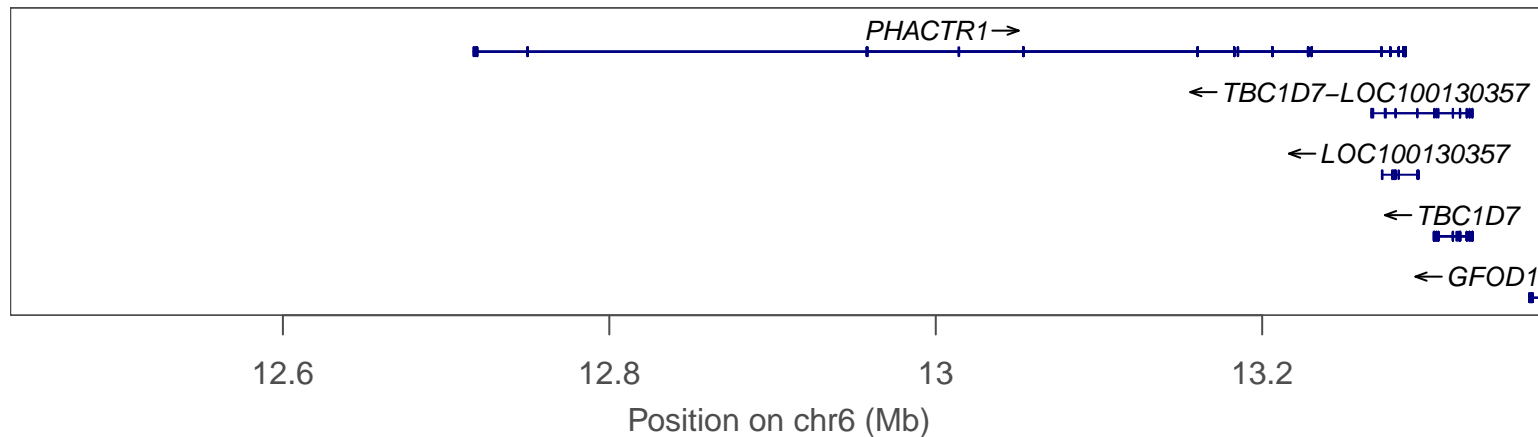
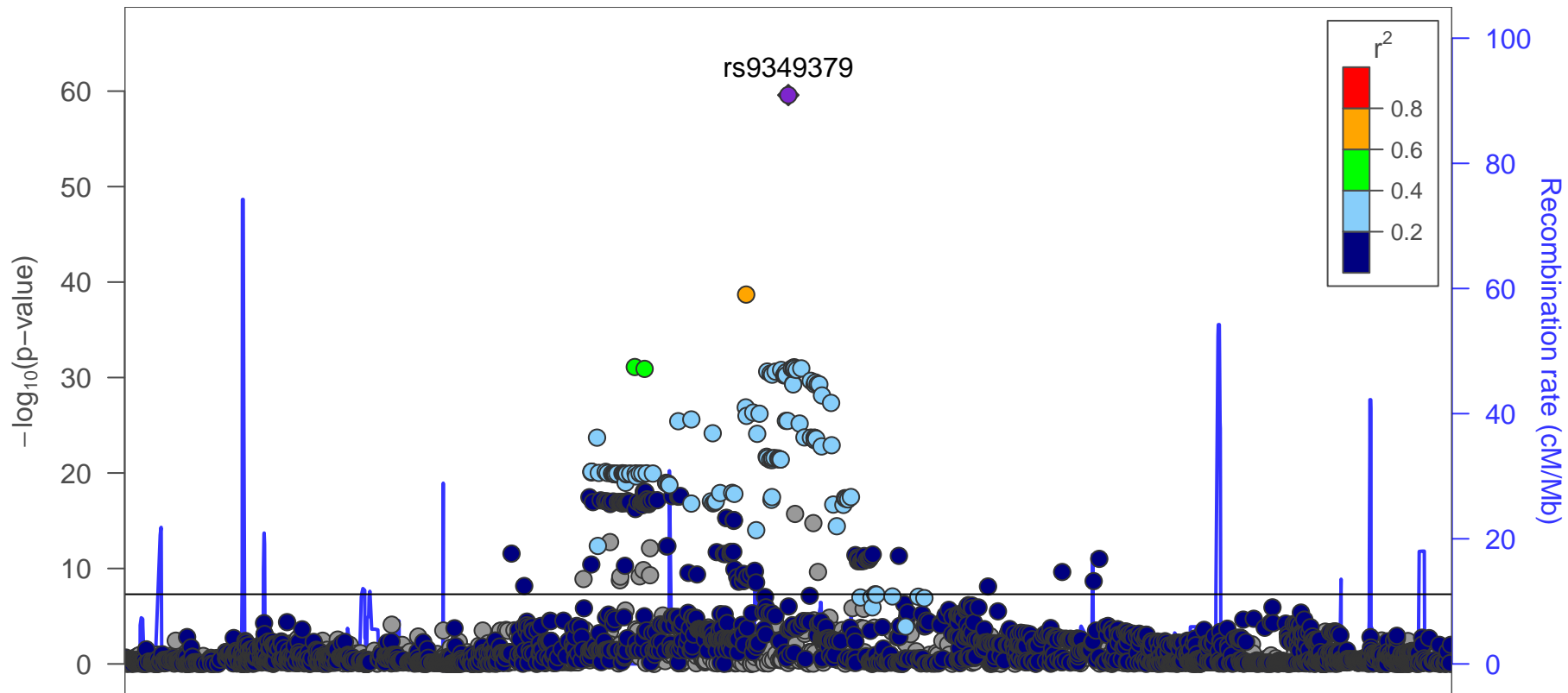
176.6

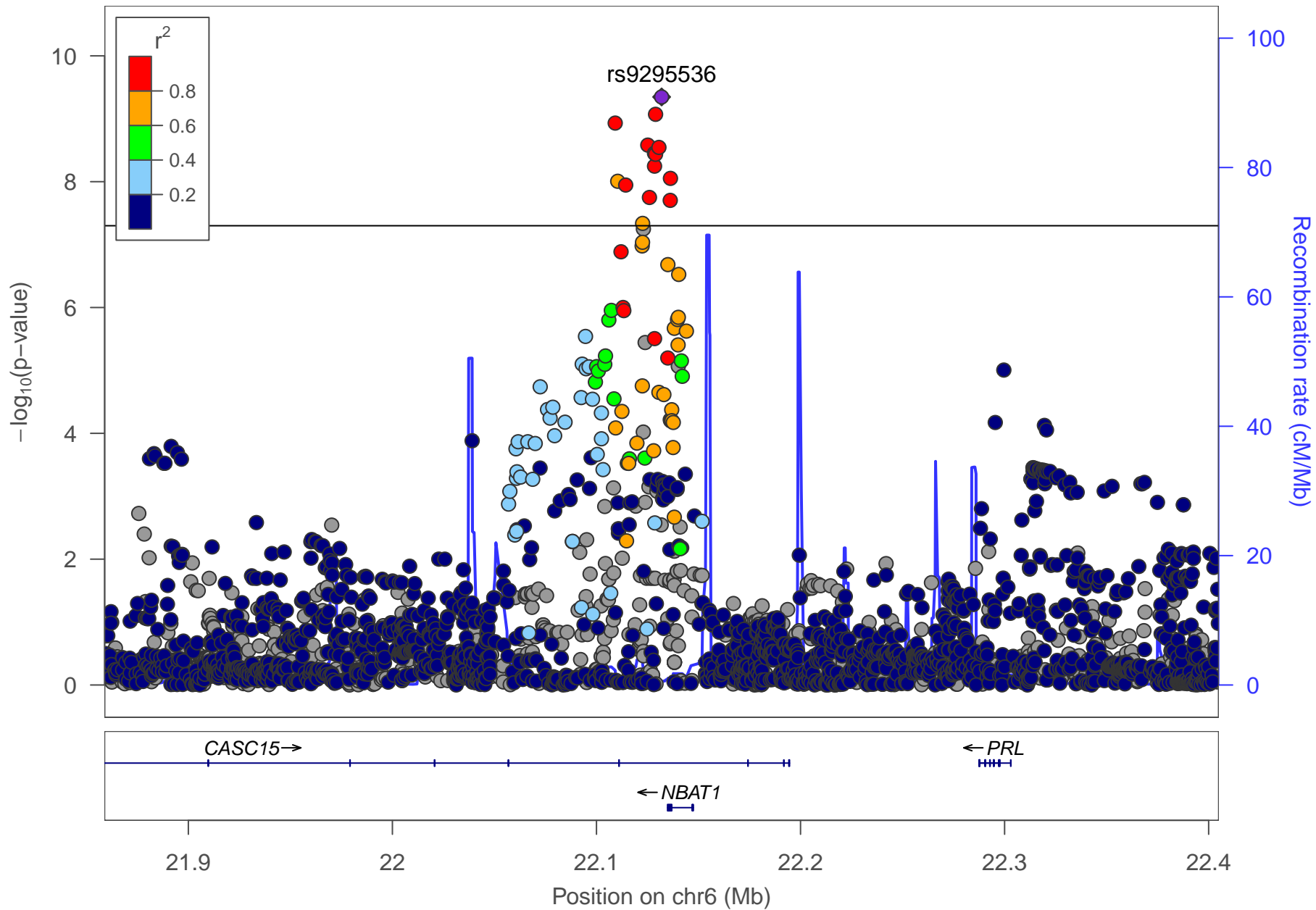
176.8

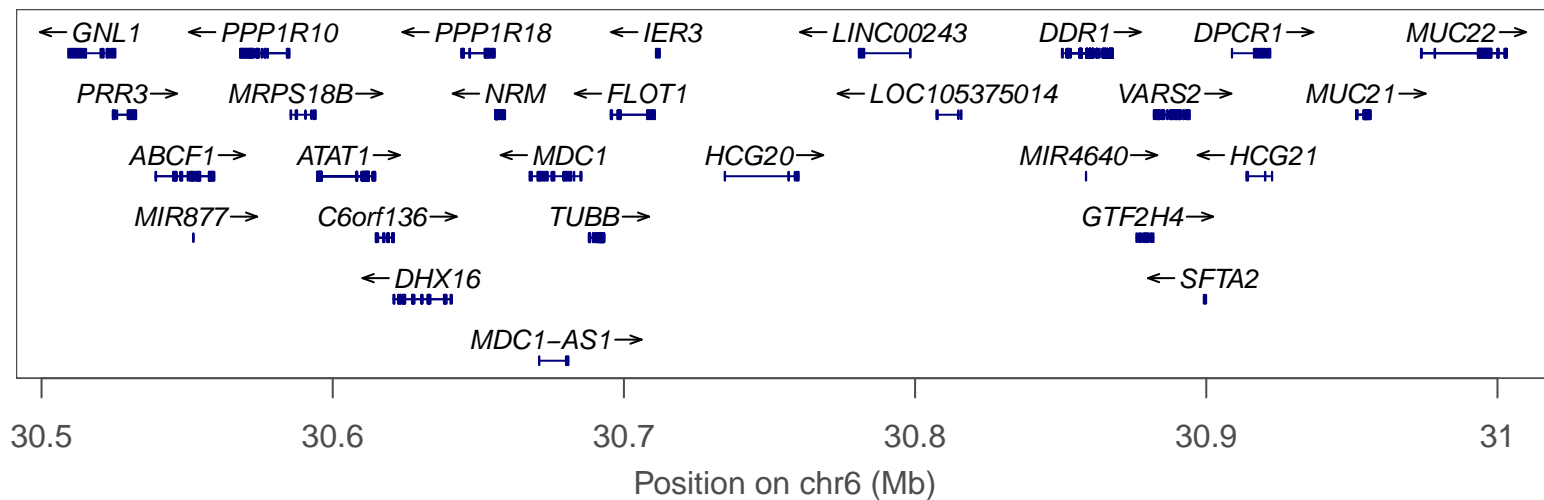
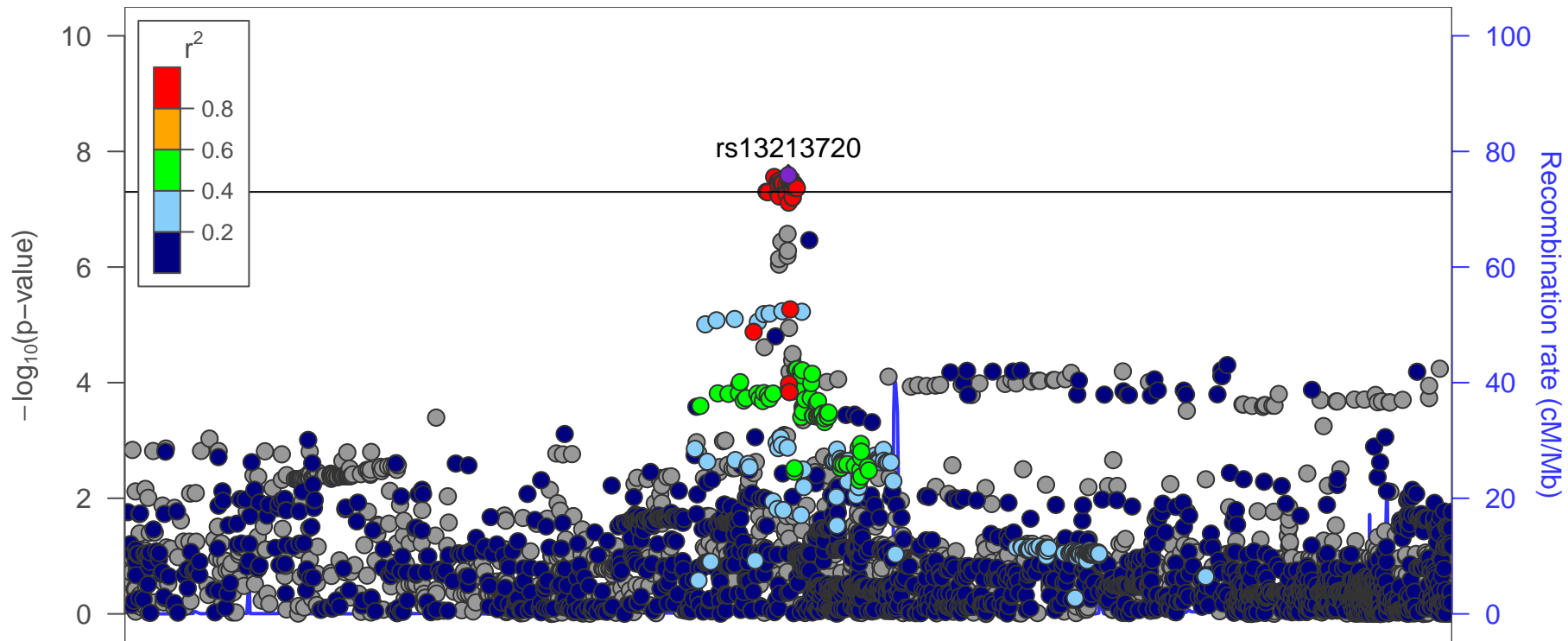
Position on chr5 (Mb)

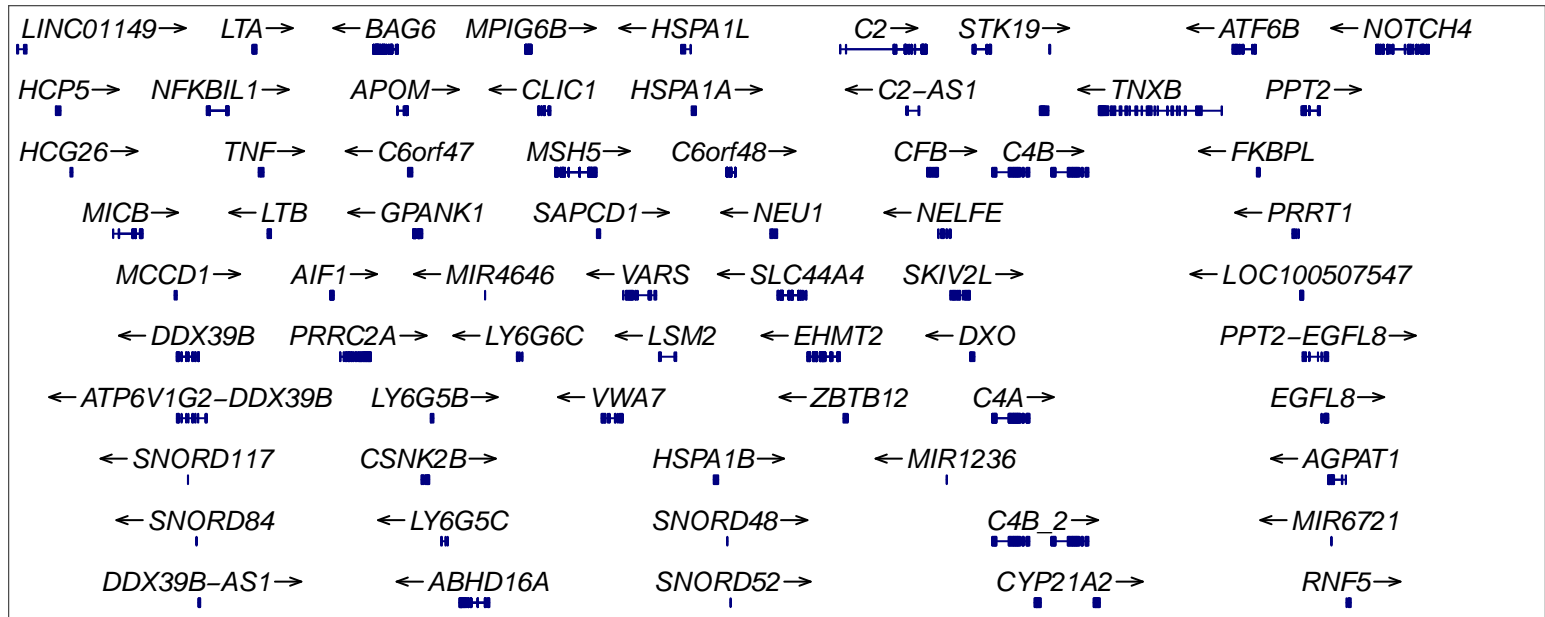
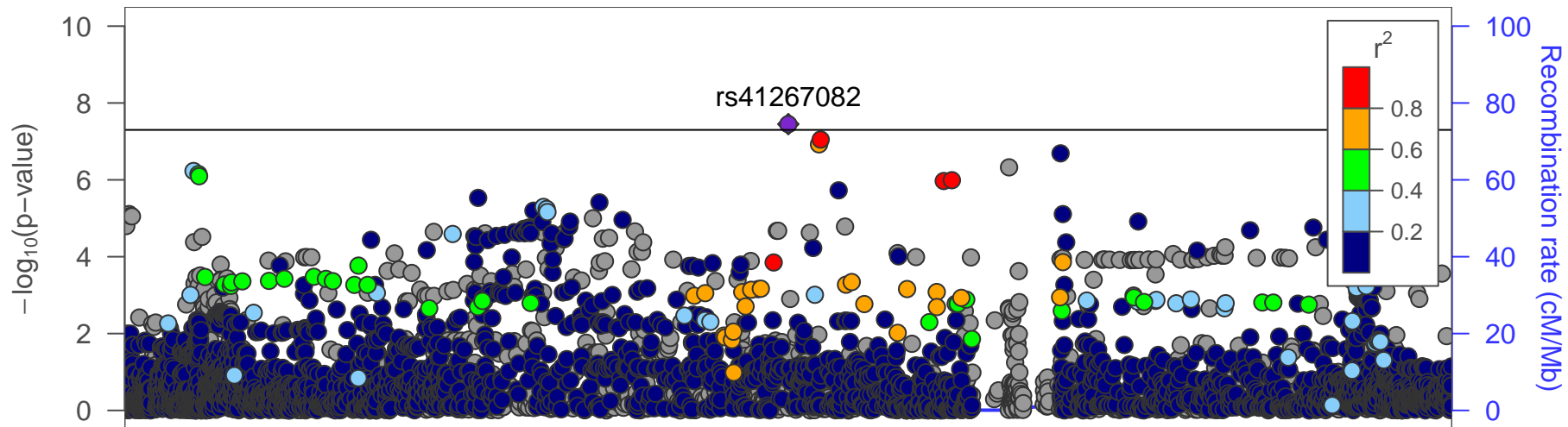












19 genes omitted

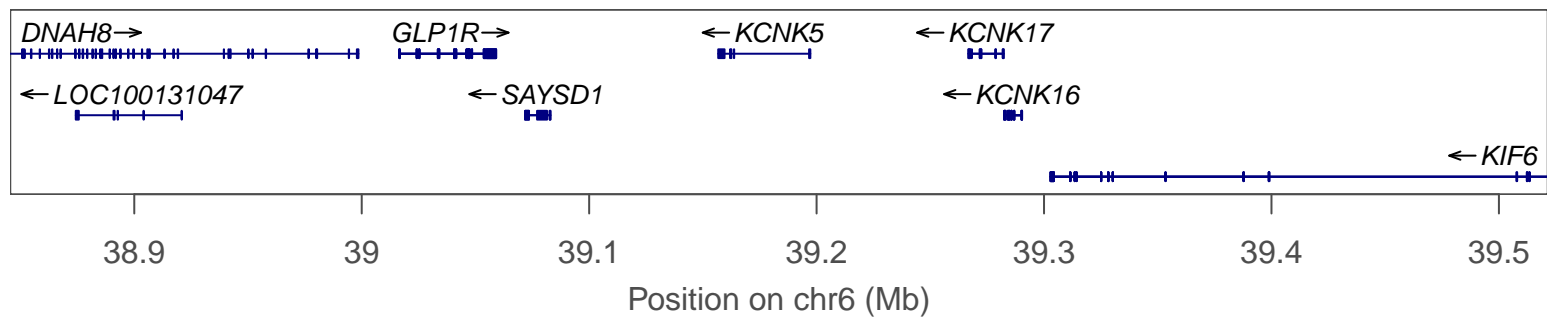
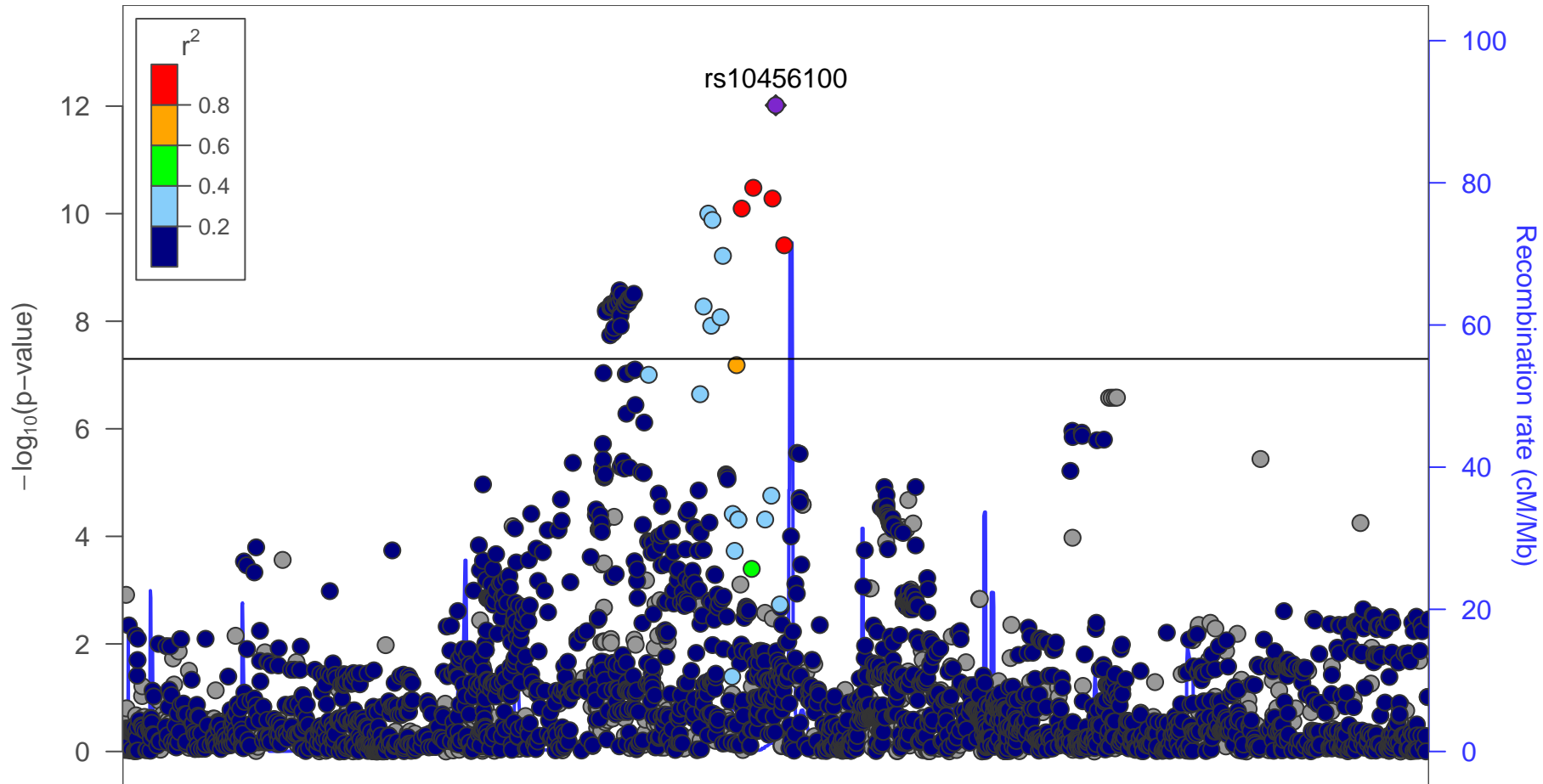
31.6

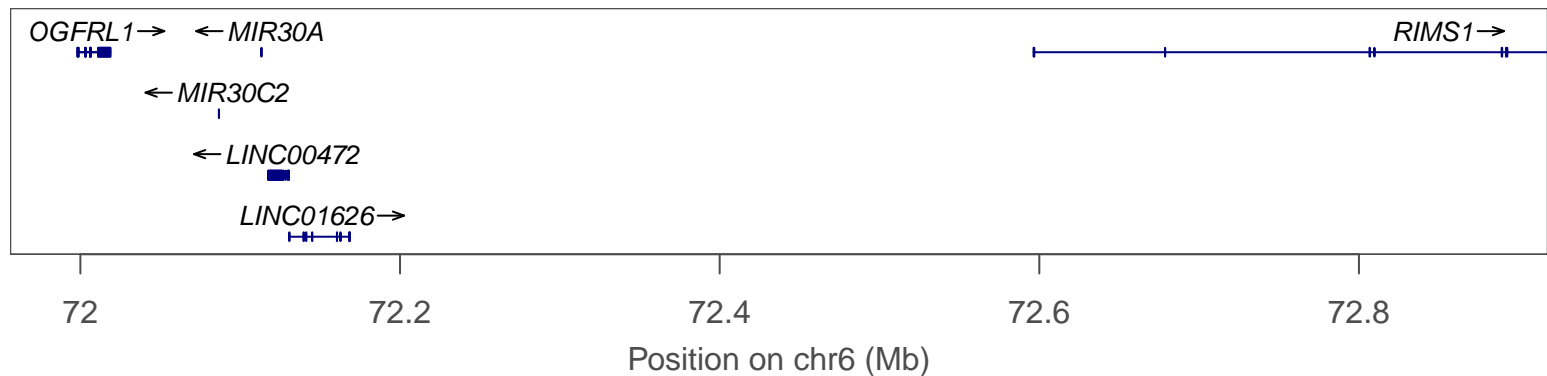
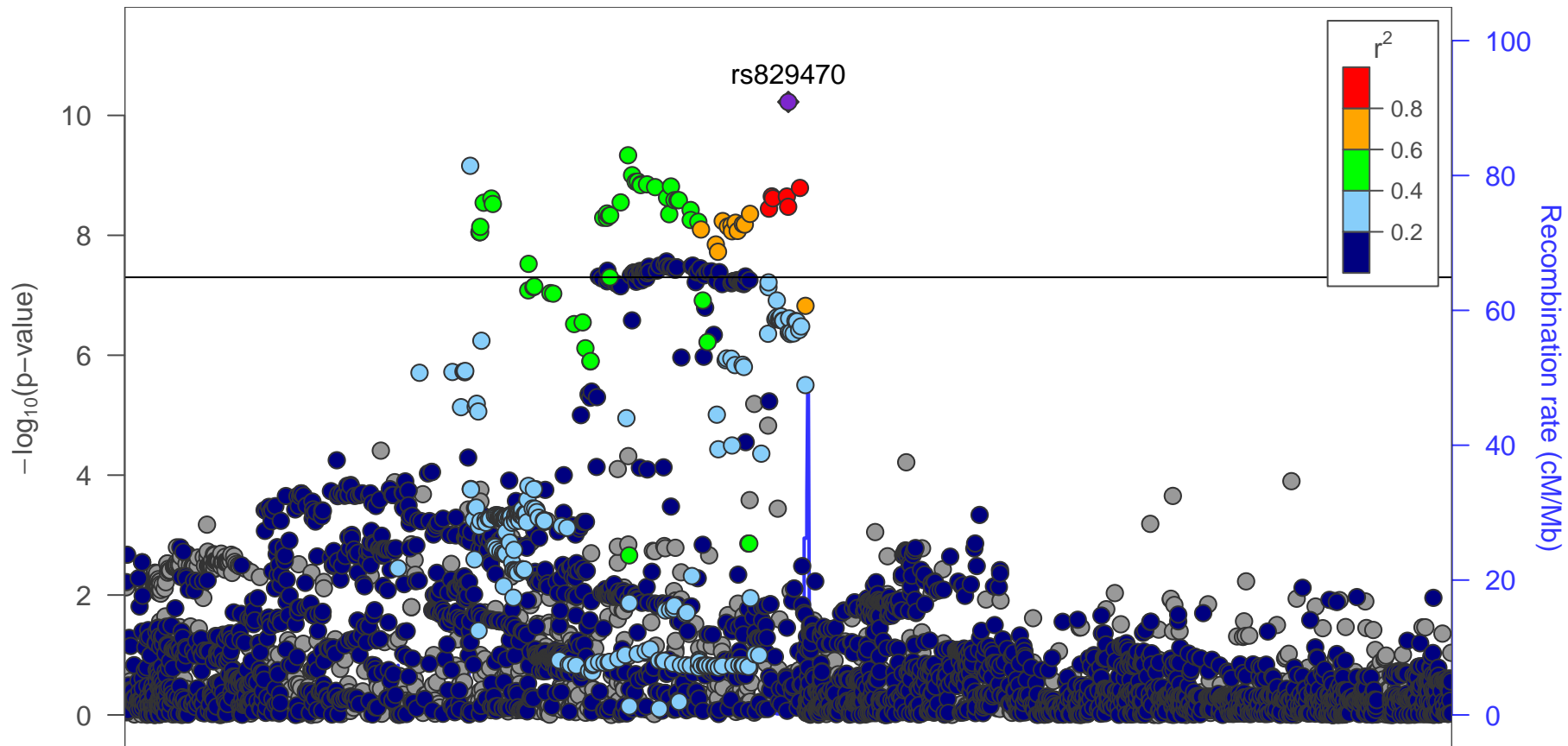
31.8

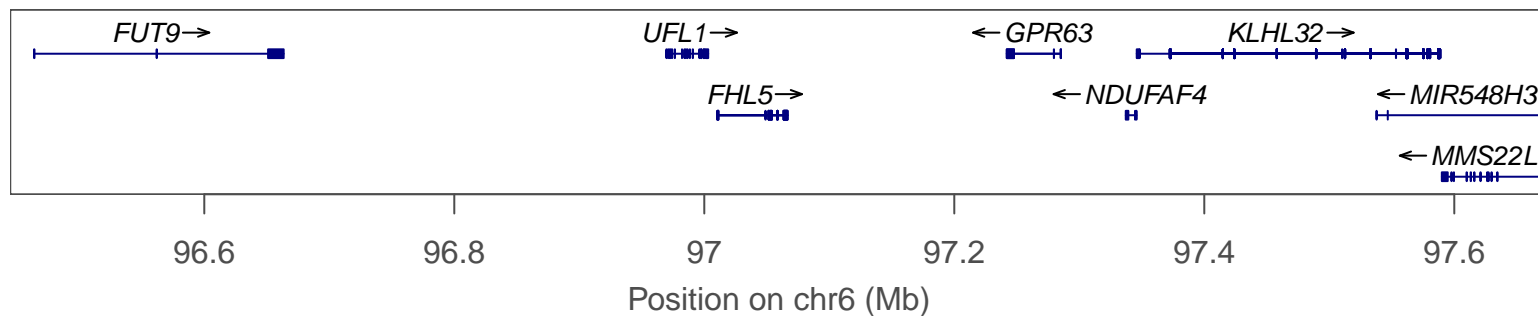
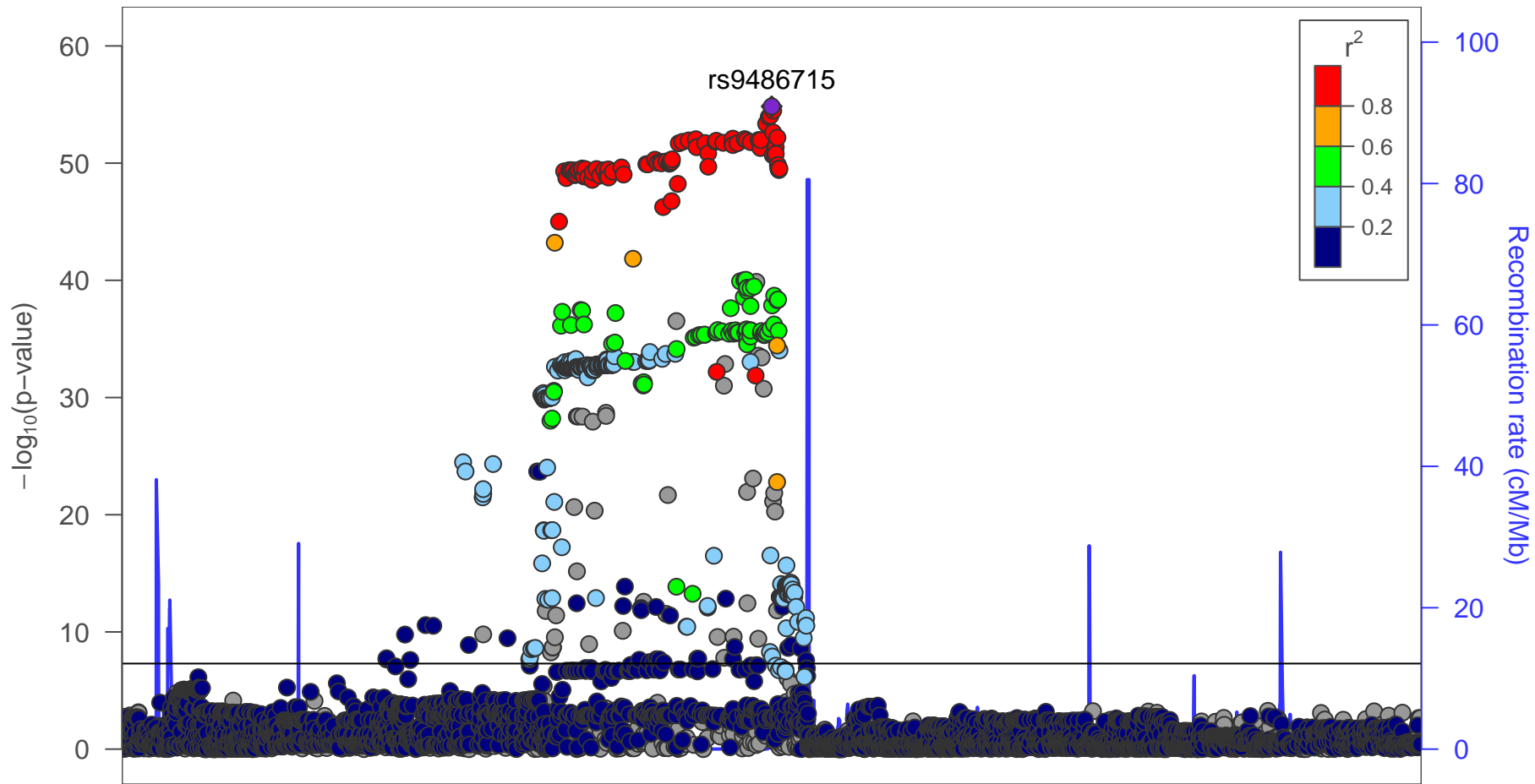
32

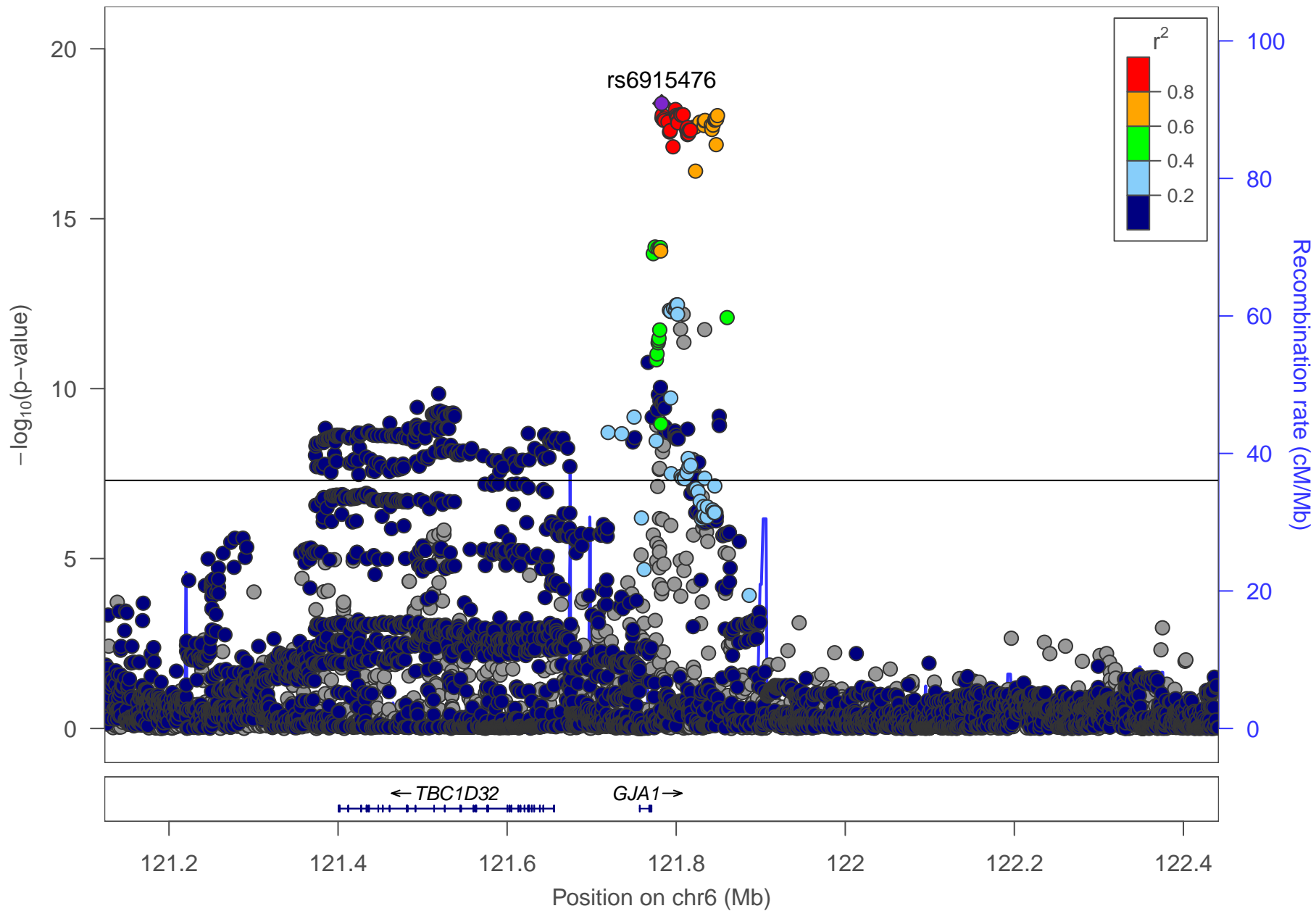
32.2

Position on chr6 (Mb)

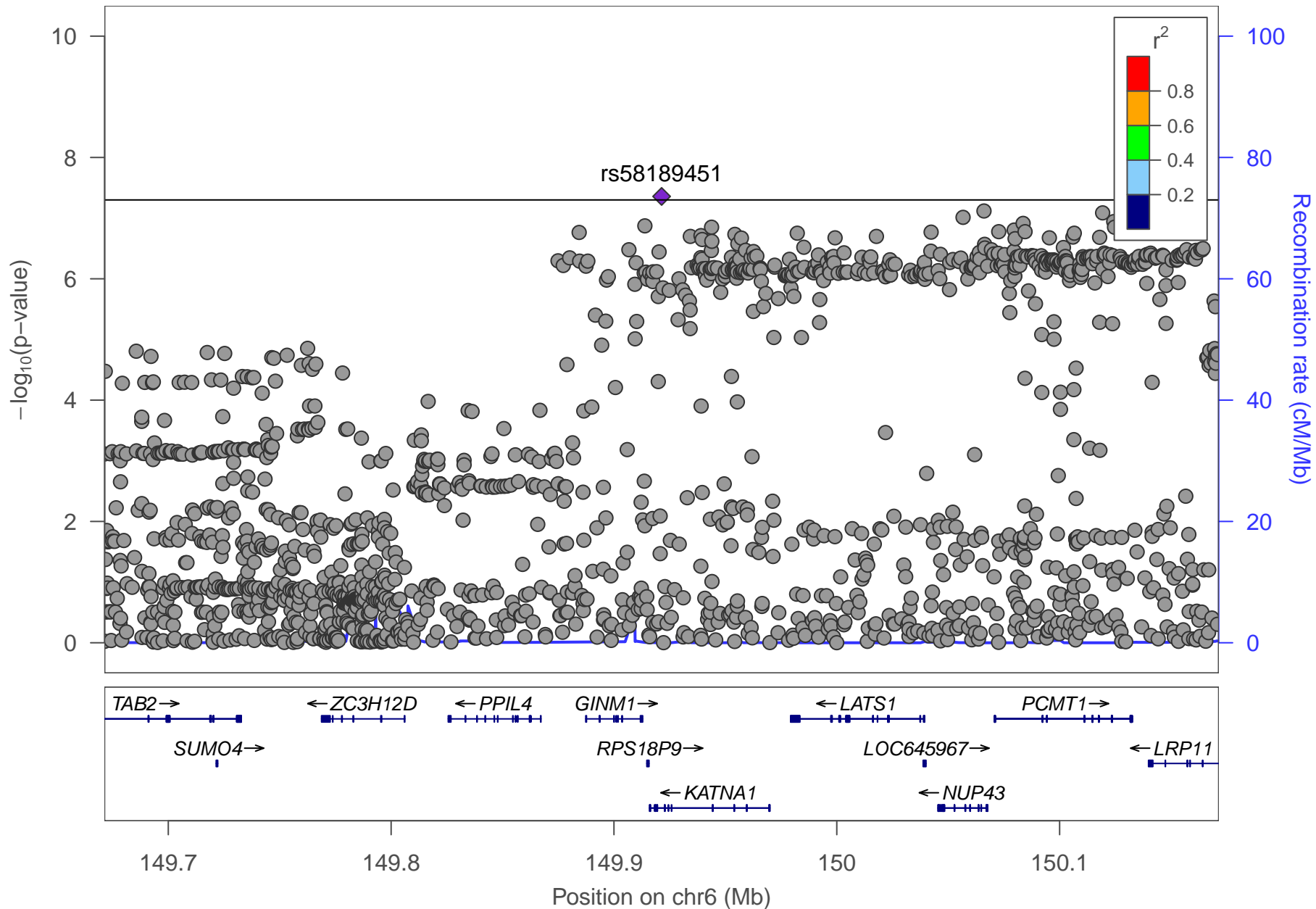


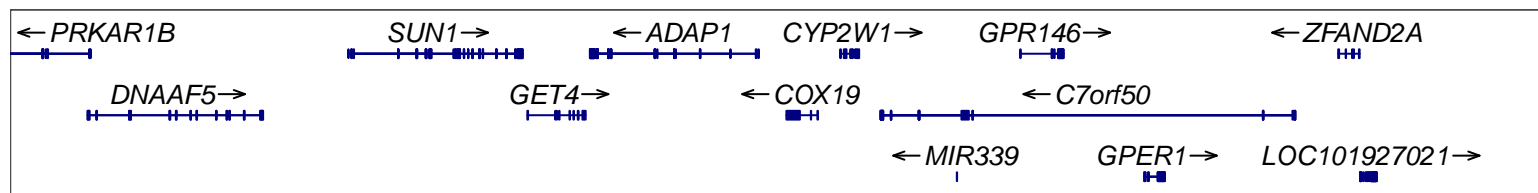
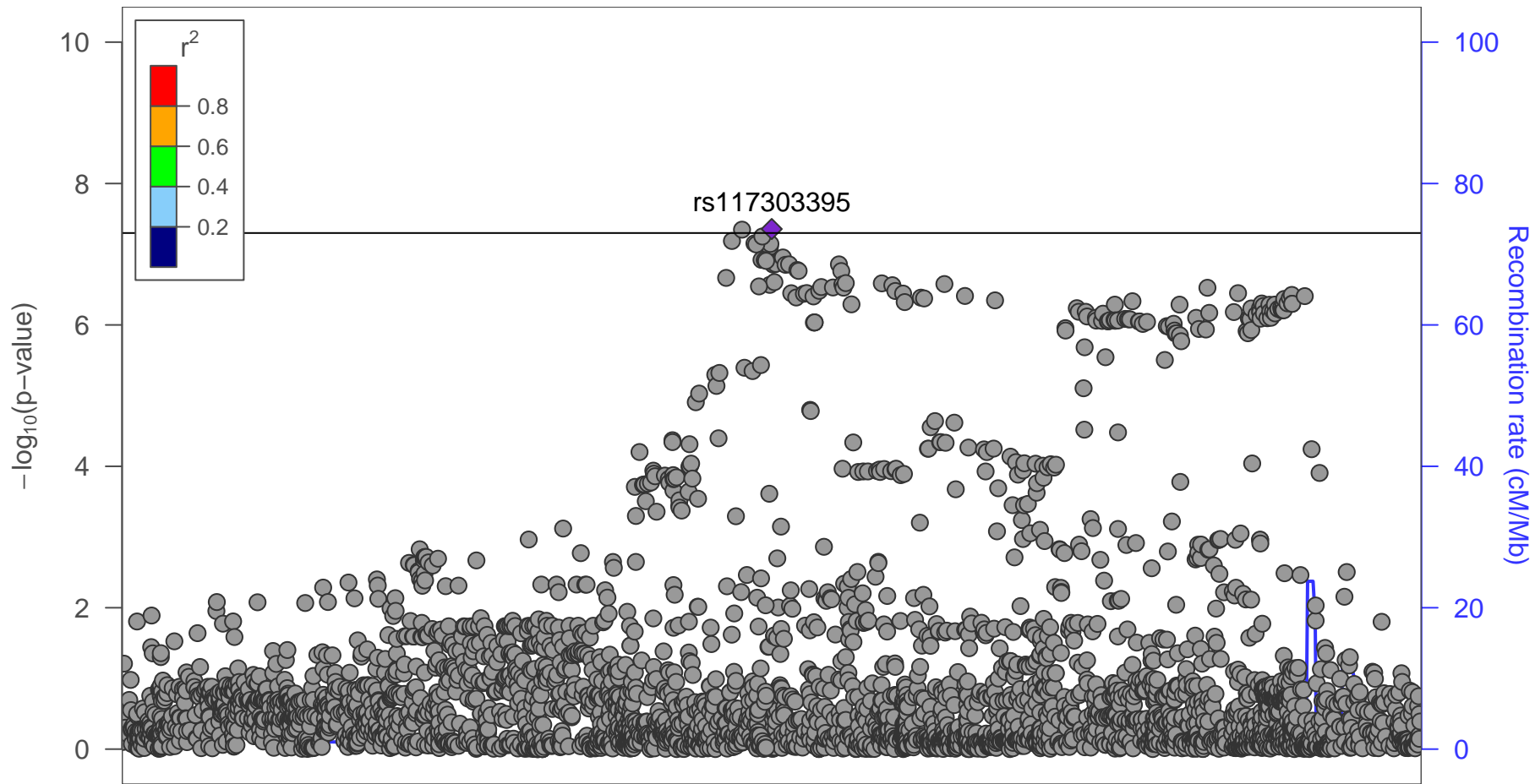












0.8                      0.9                      1                      1.1                      1.2

Position on chr7 (Mb)

