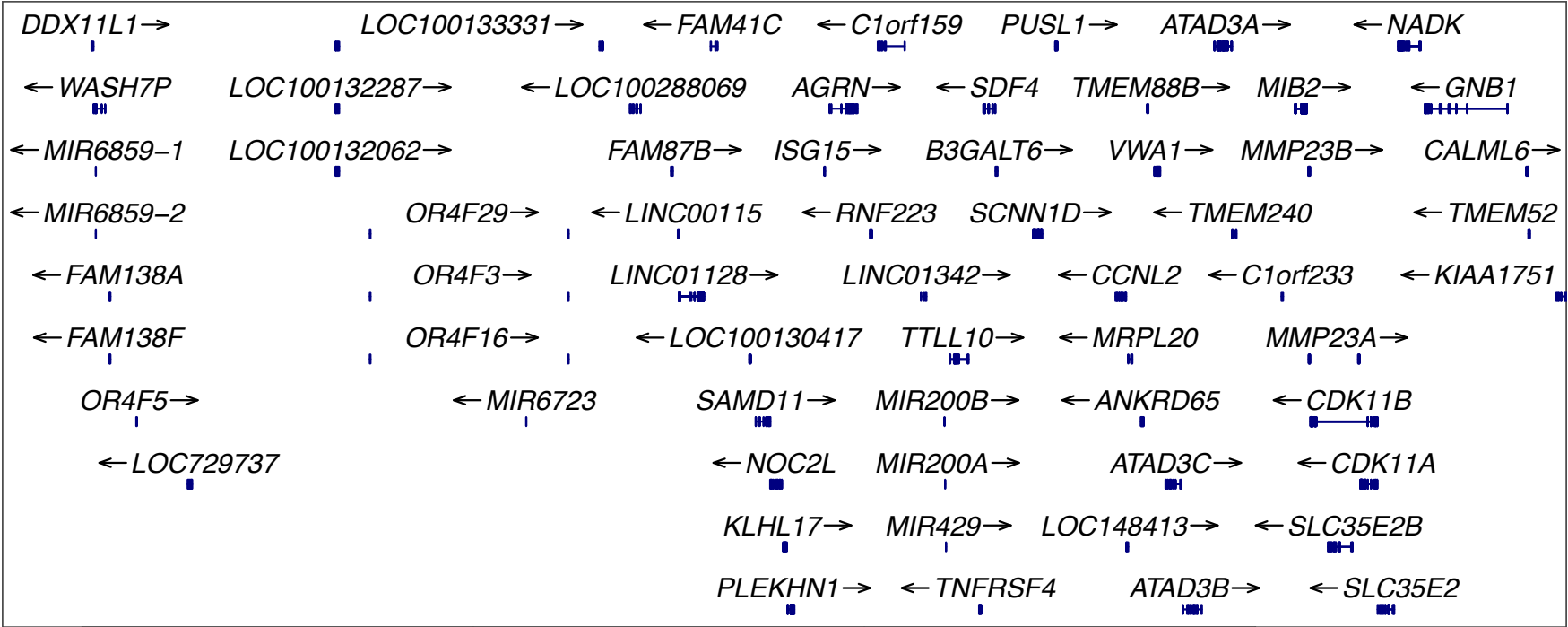
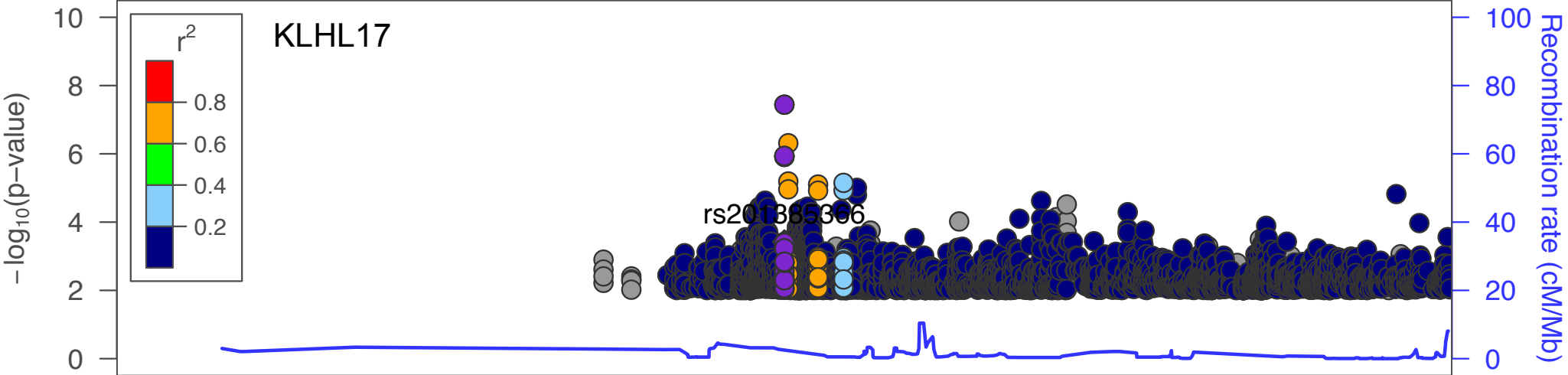


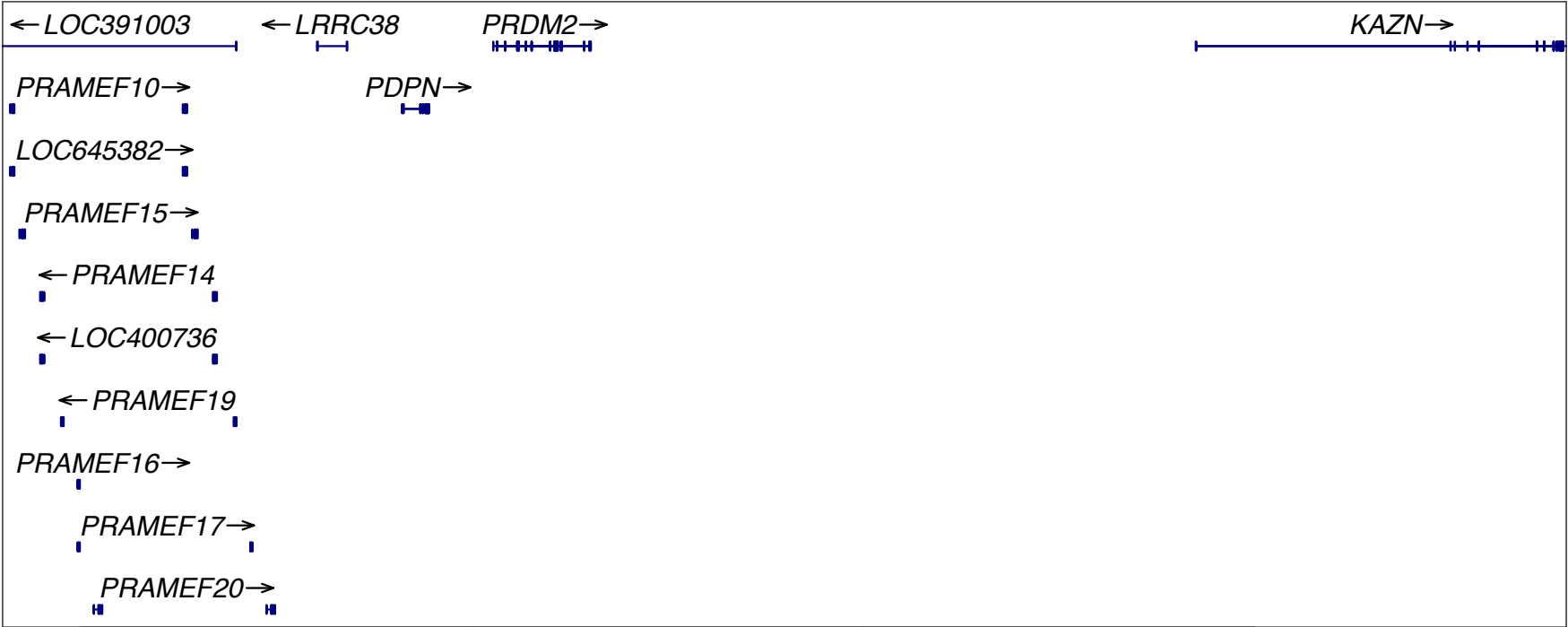
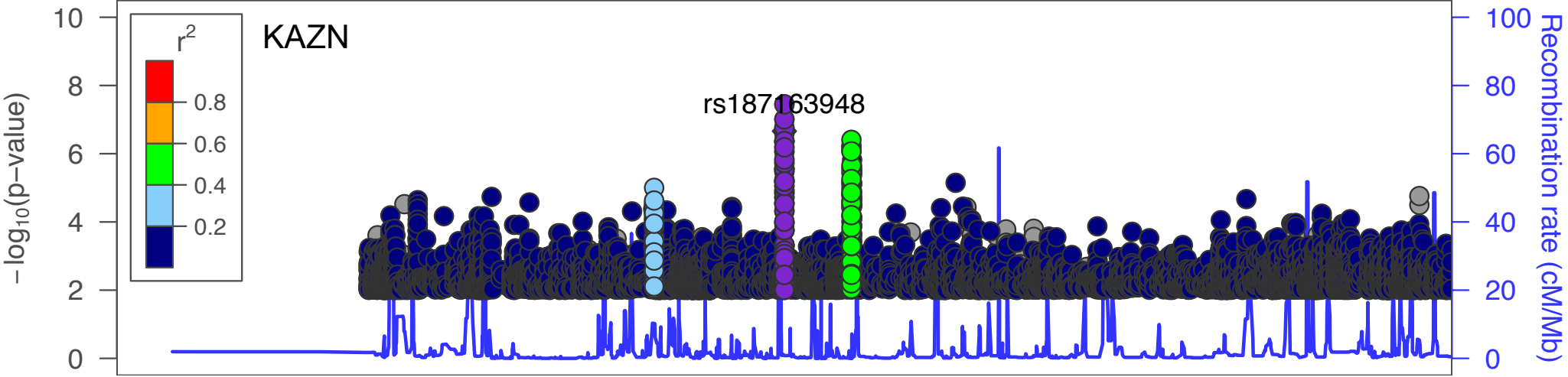
Plotted SNPs



16 genes
omitted

Position on chr1 (Mb)

Plotted SNPs



2 genes
omitted

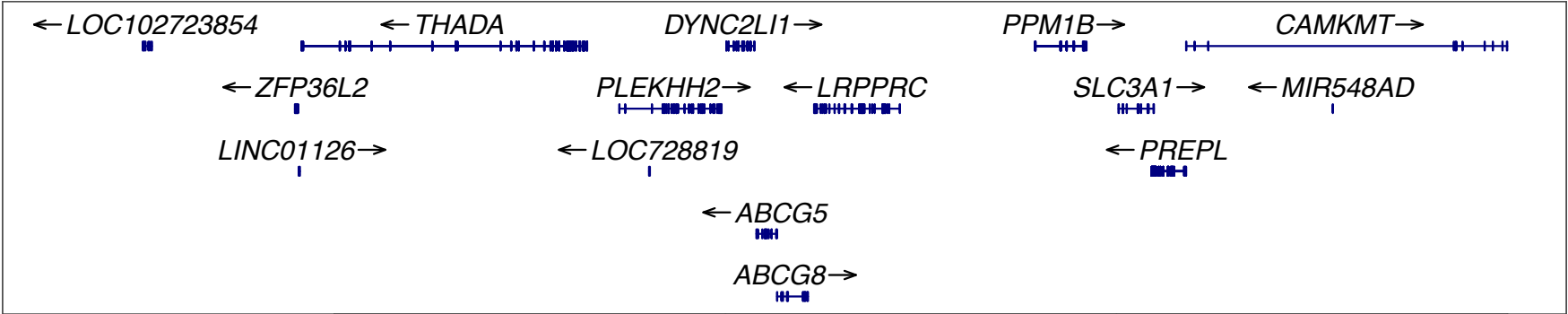
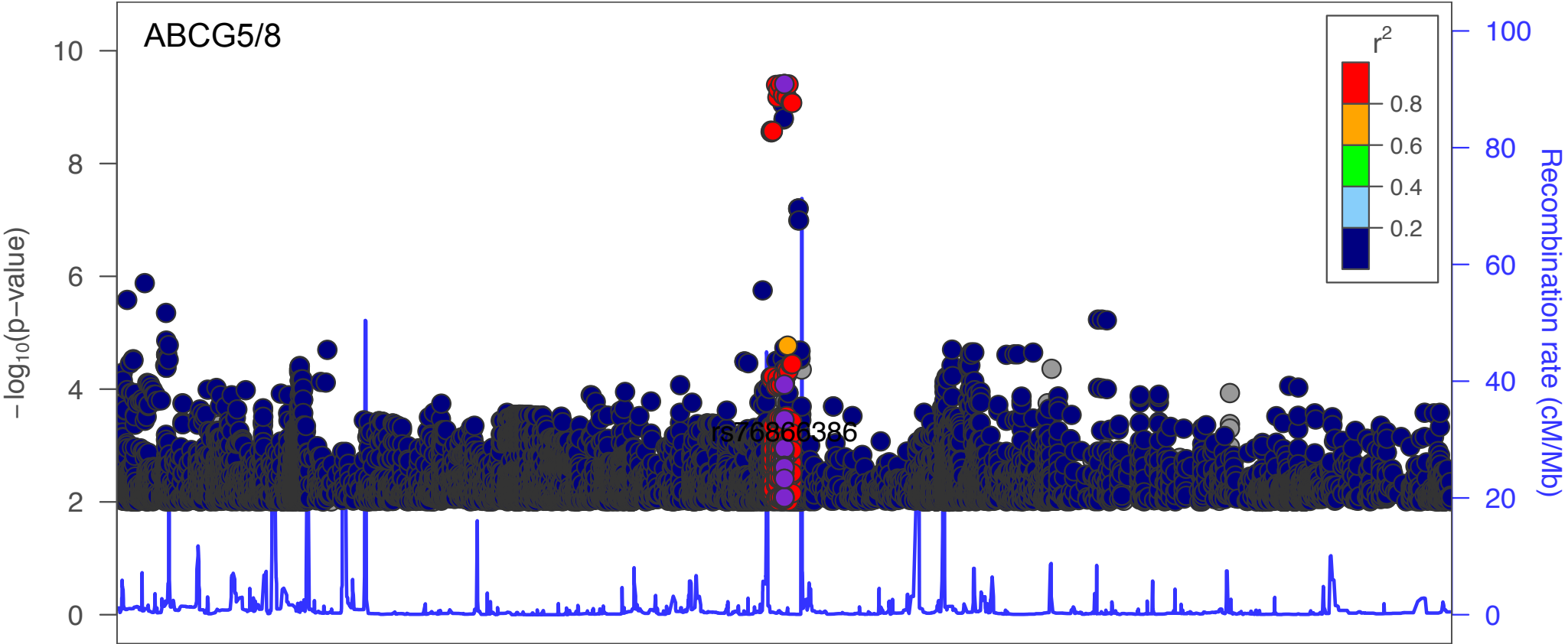
13.5

14

14.5

15

Position on chr1 (Mb)



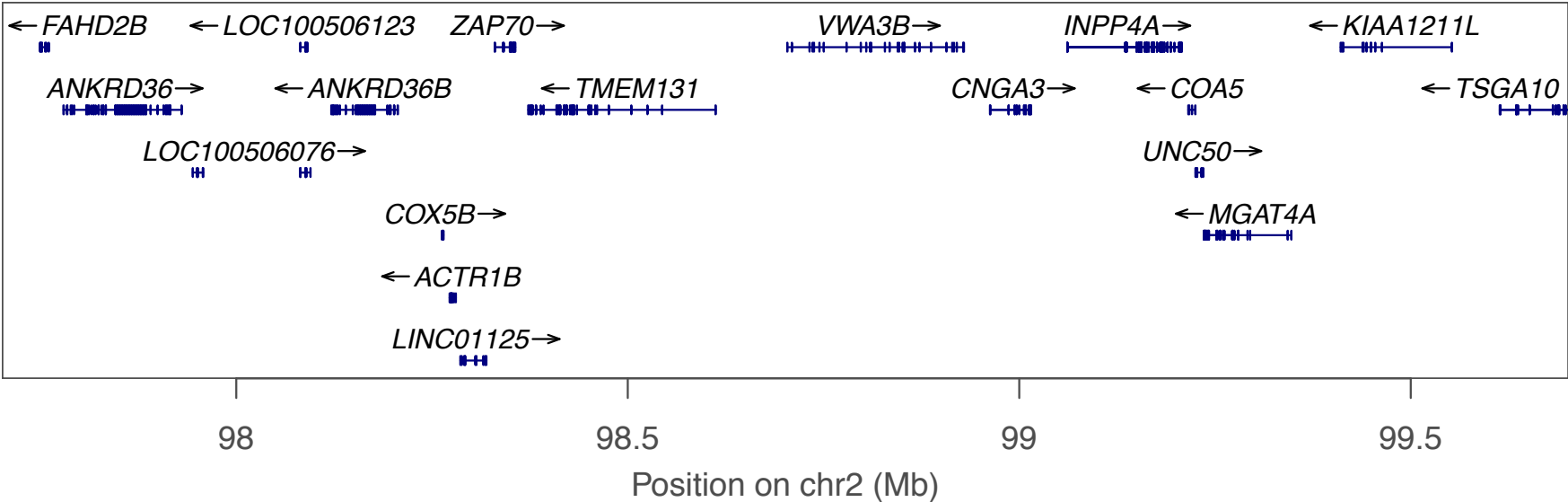
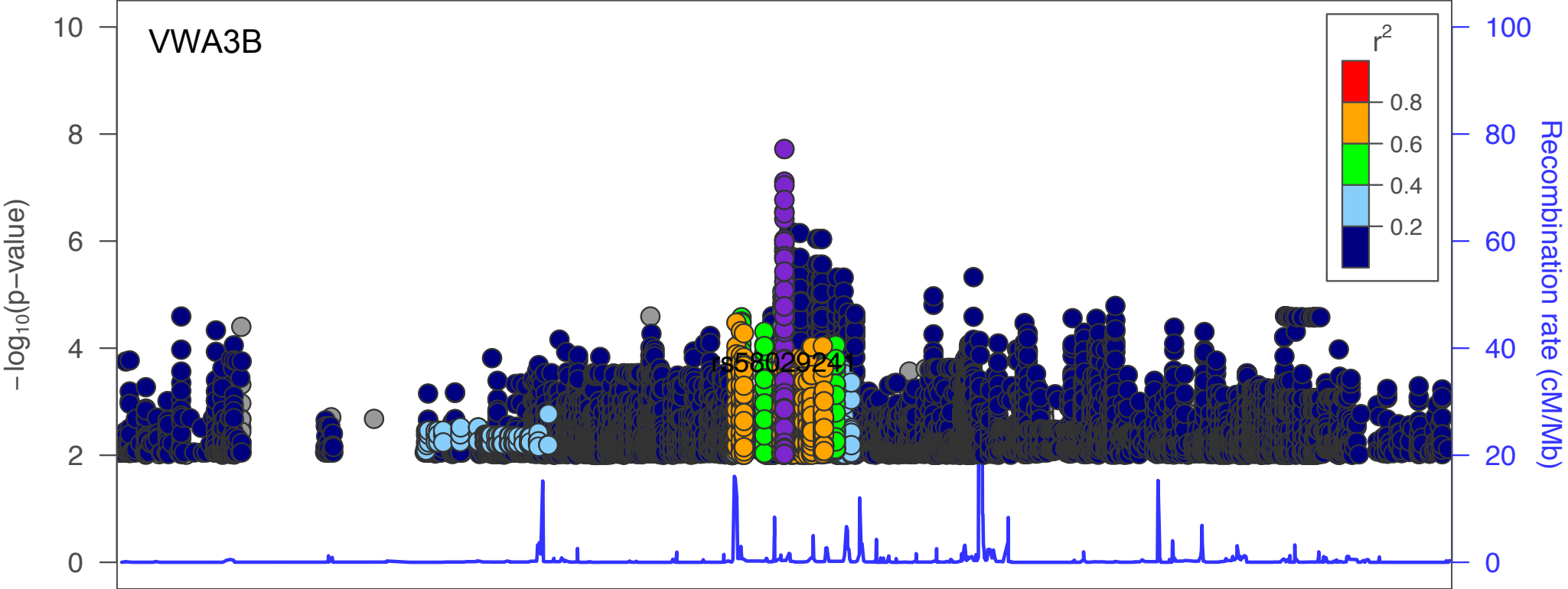
43.5

44

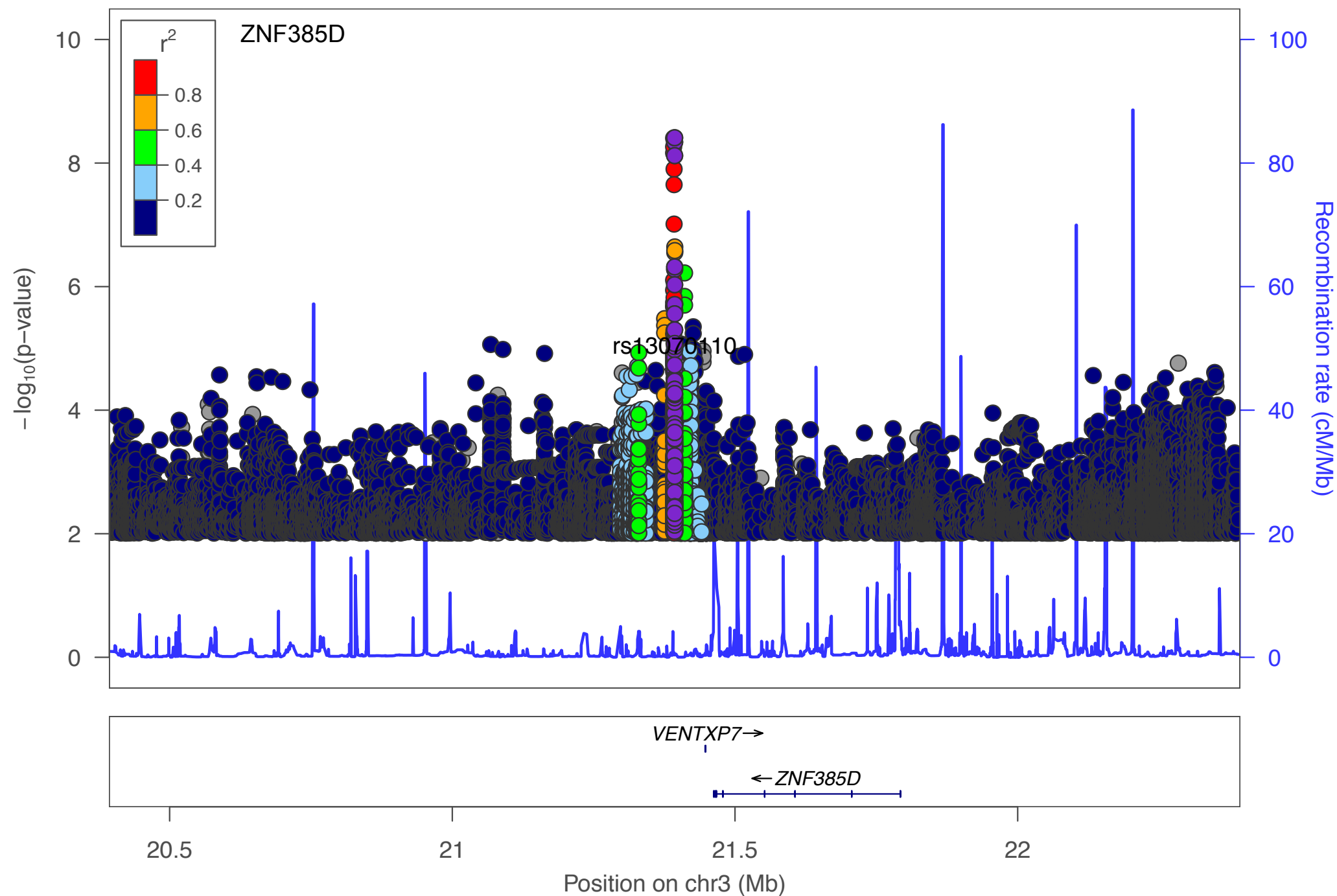
44.5

45

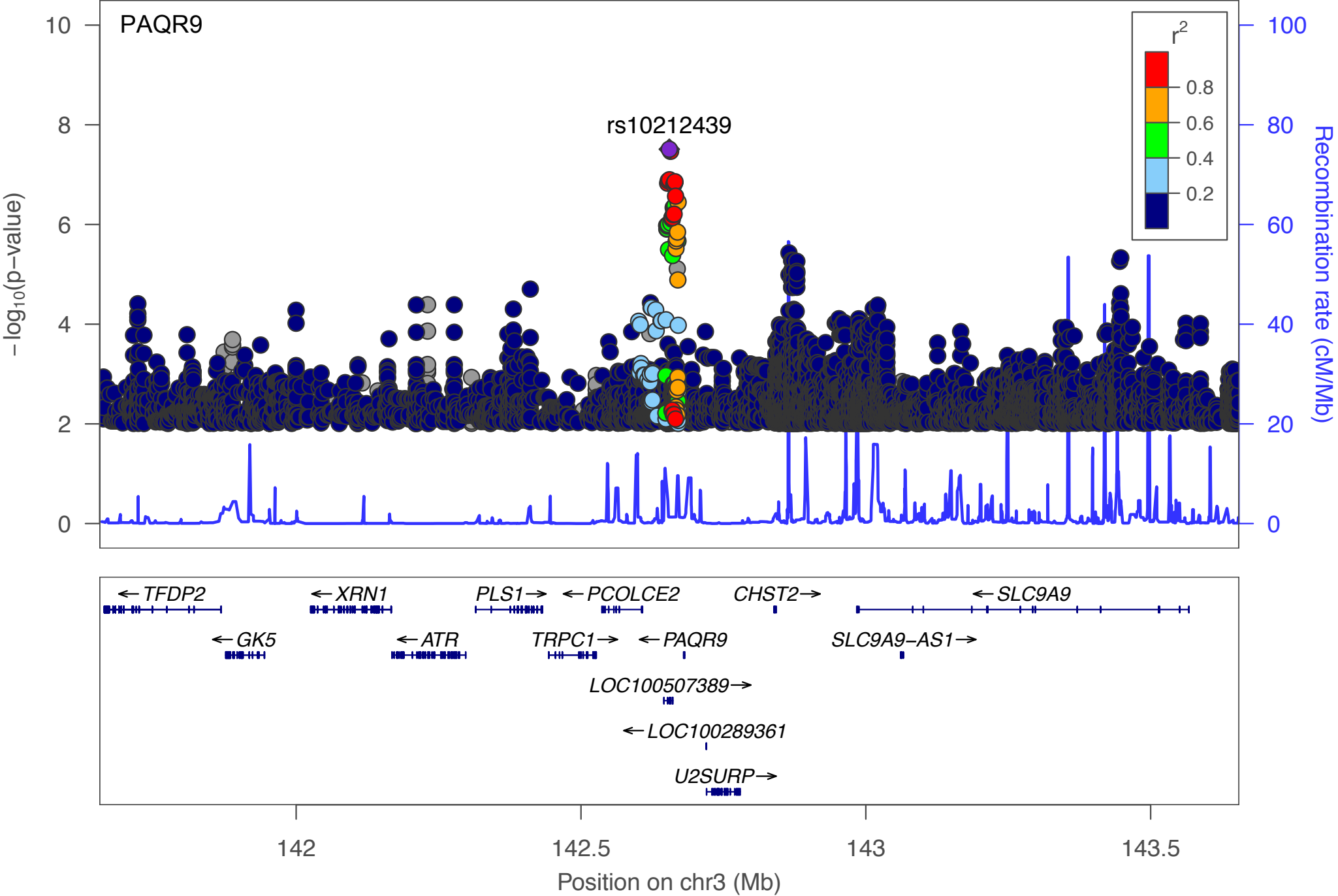
Position on chr2 (Mb)

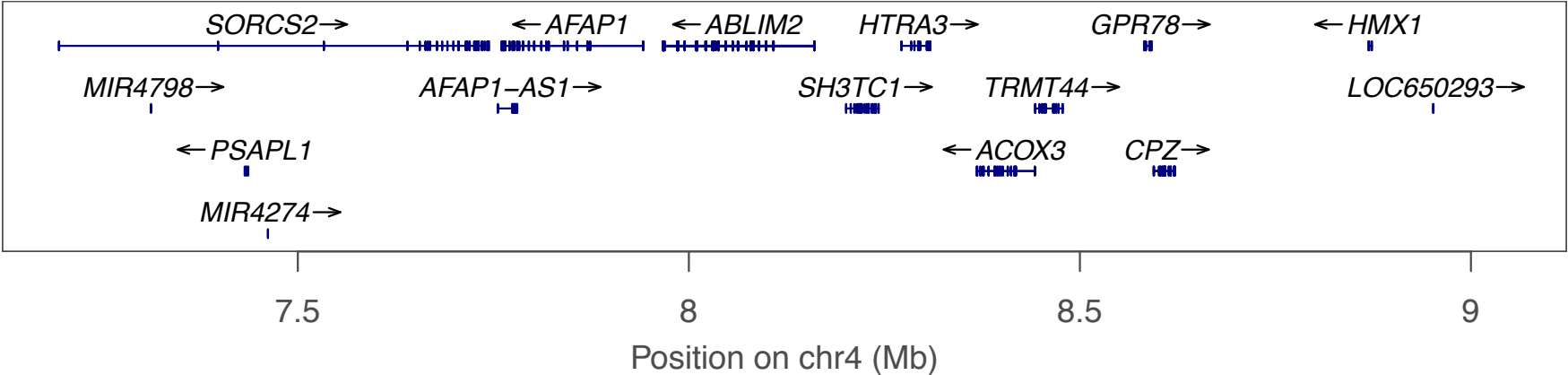
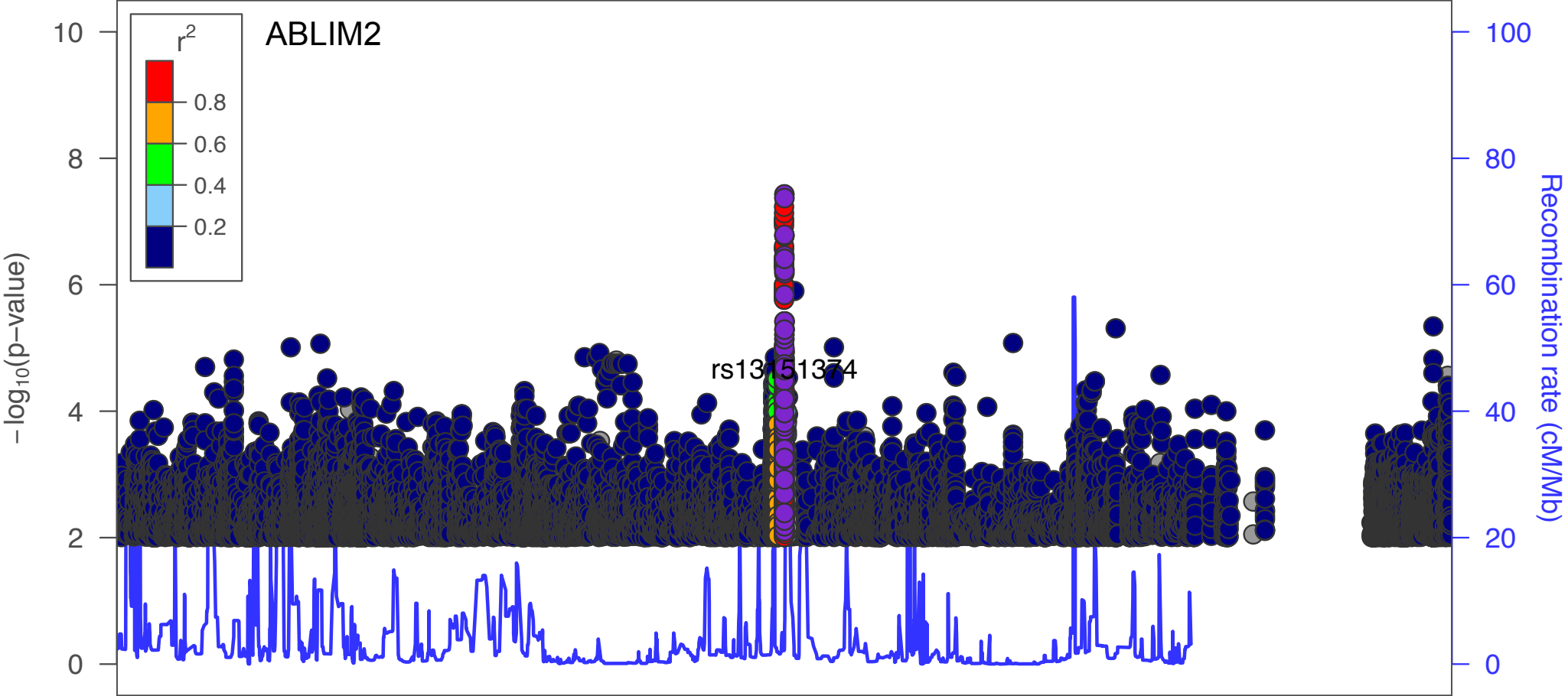


Plotted SNPs

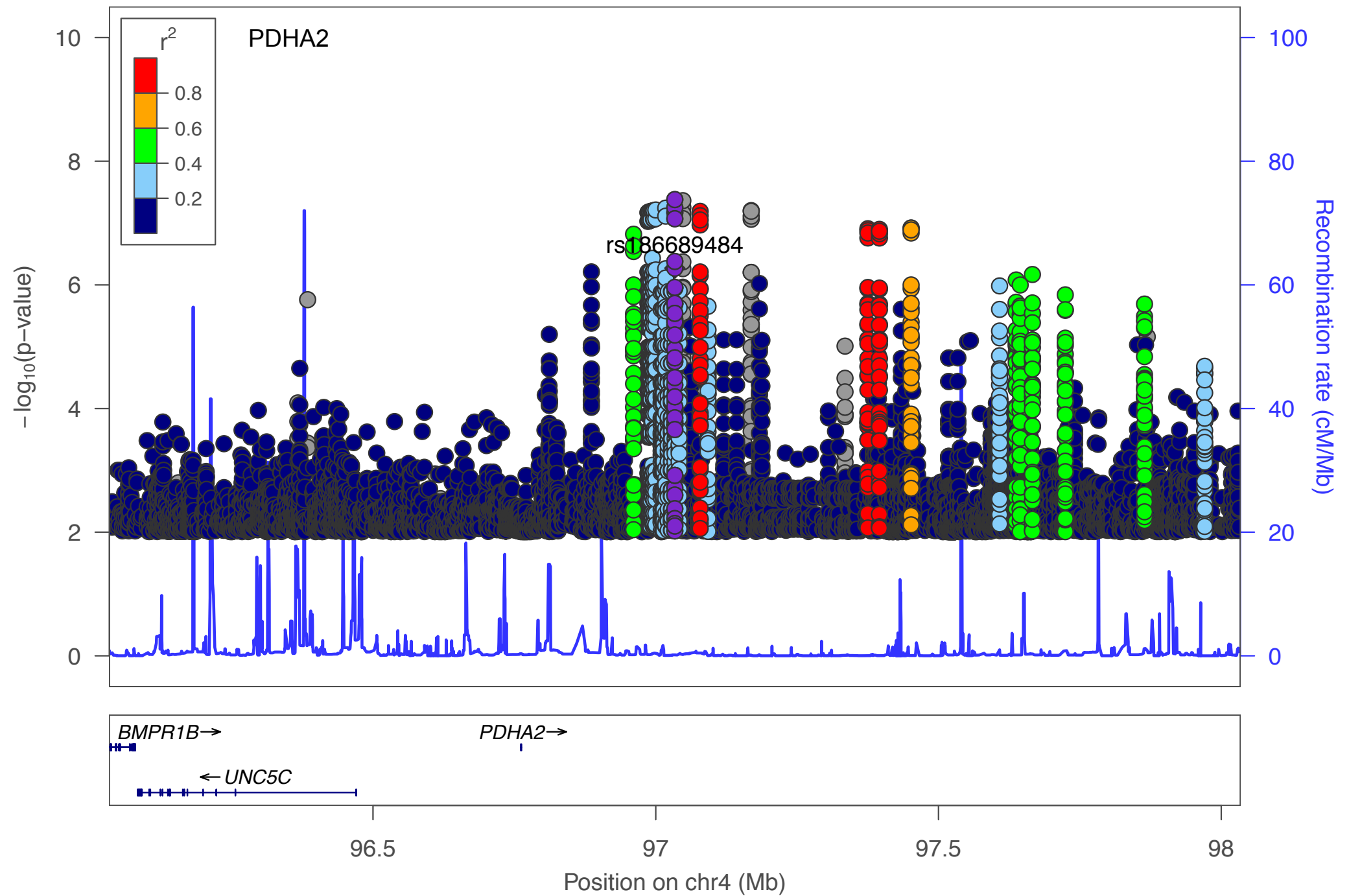


Plotted SNPs

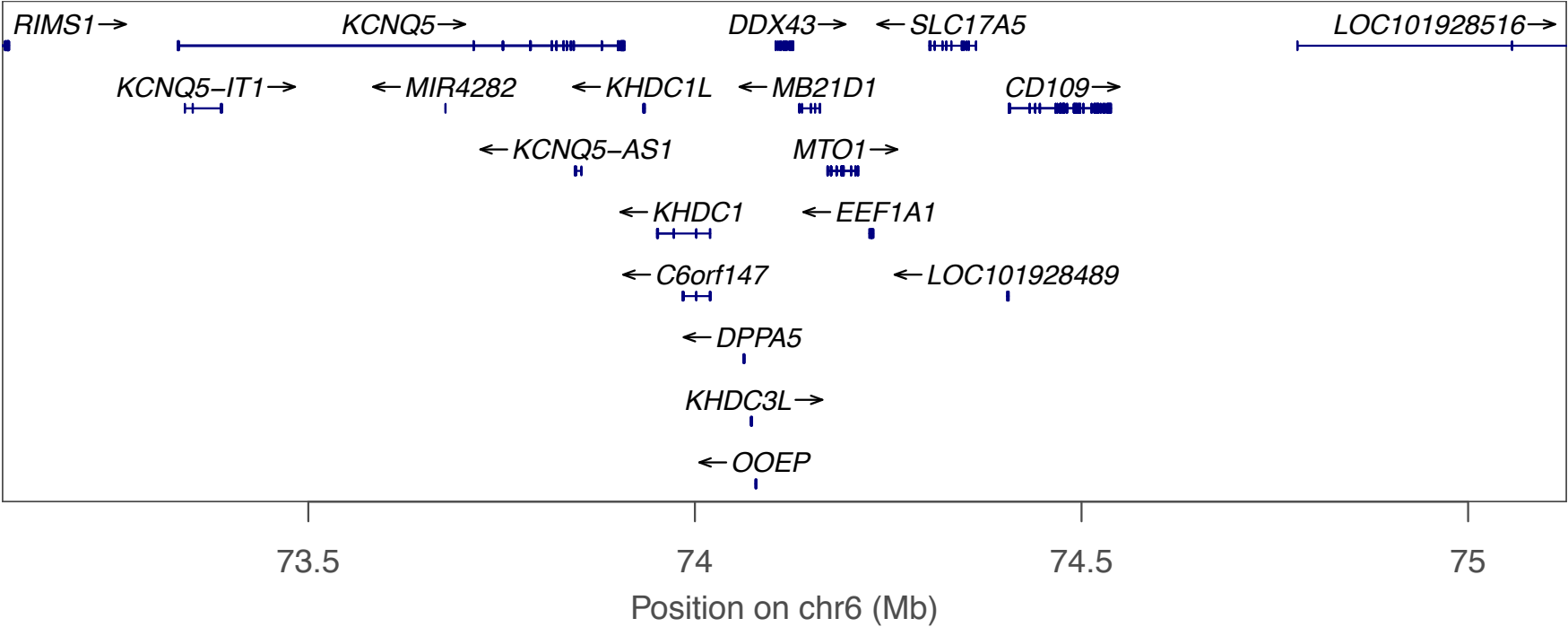
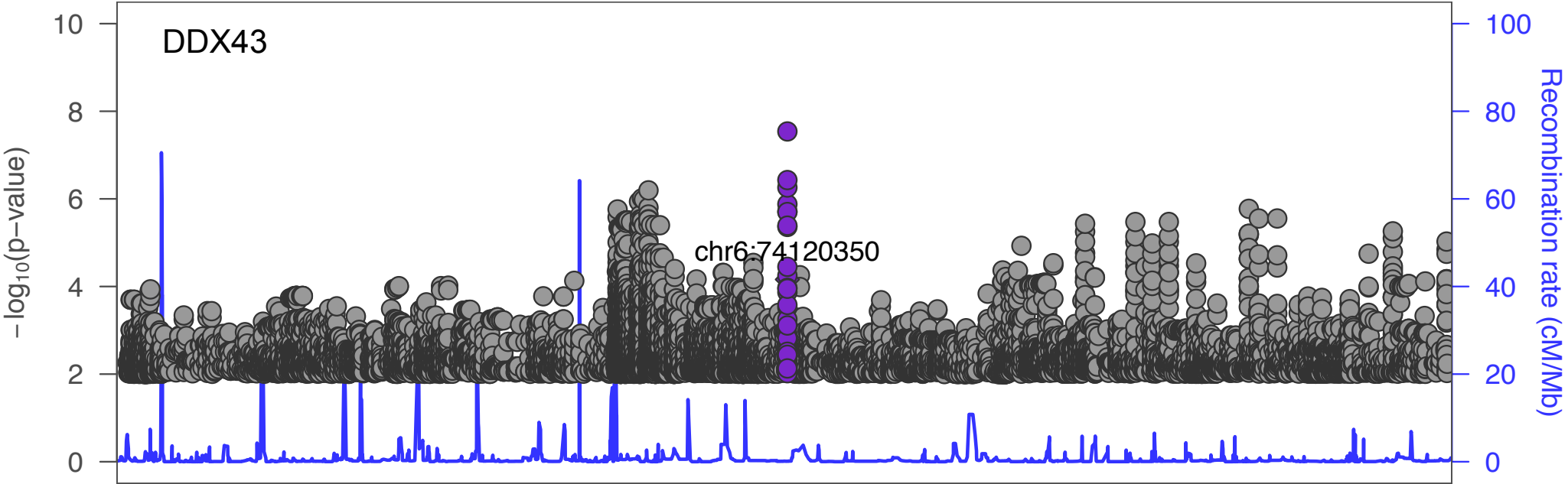




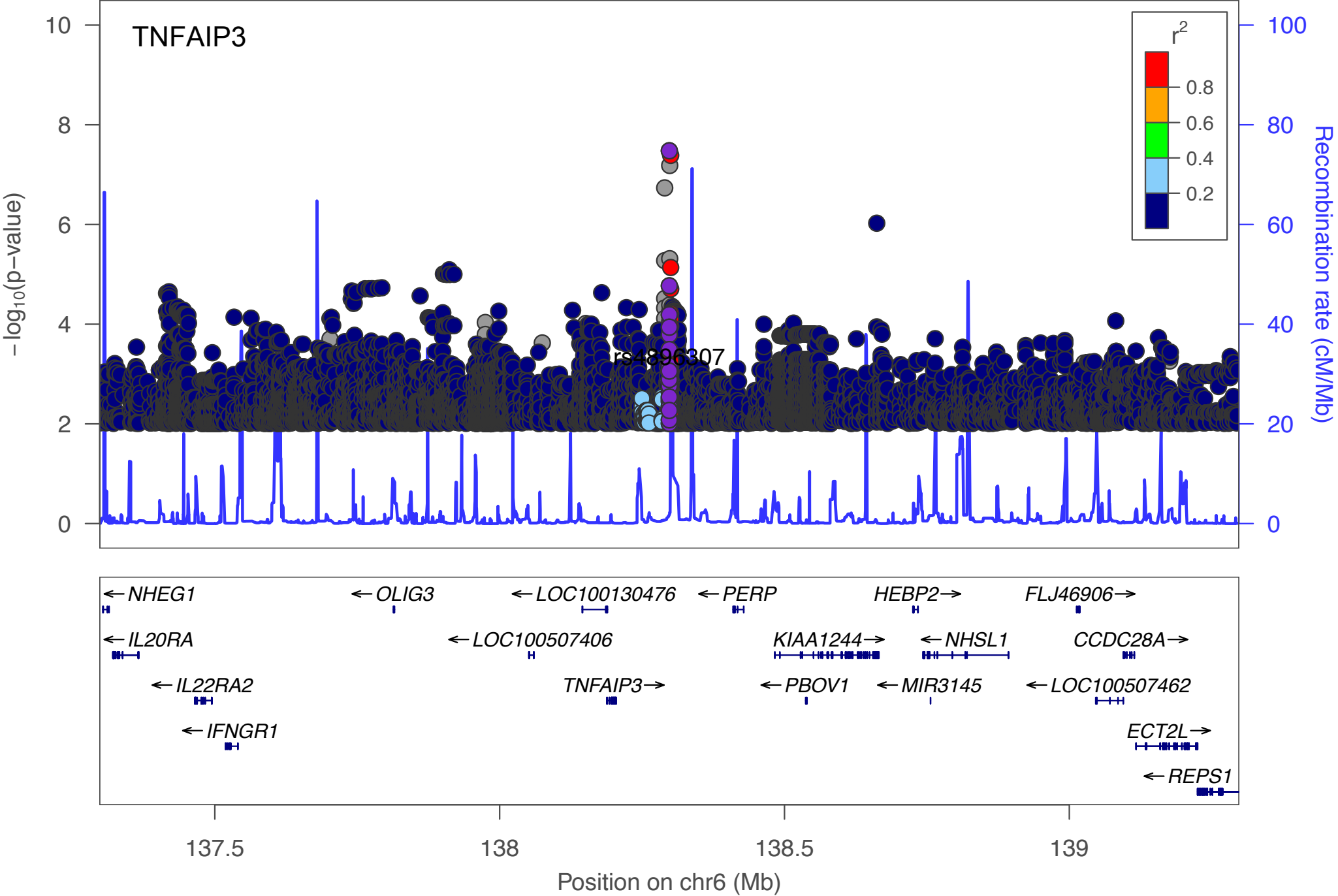
Plotted SNPs



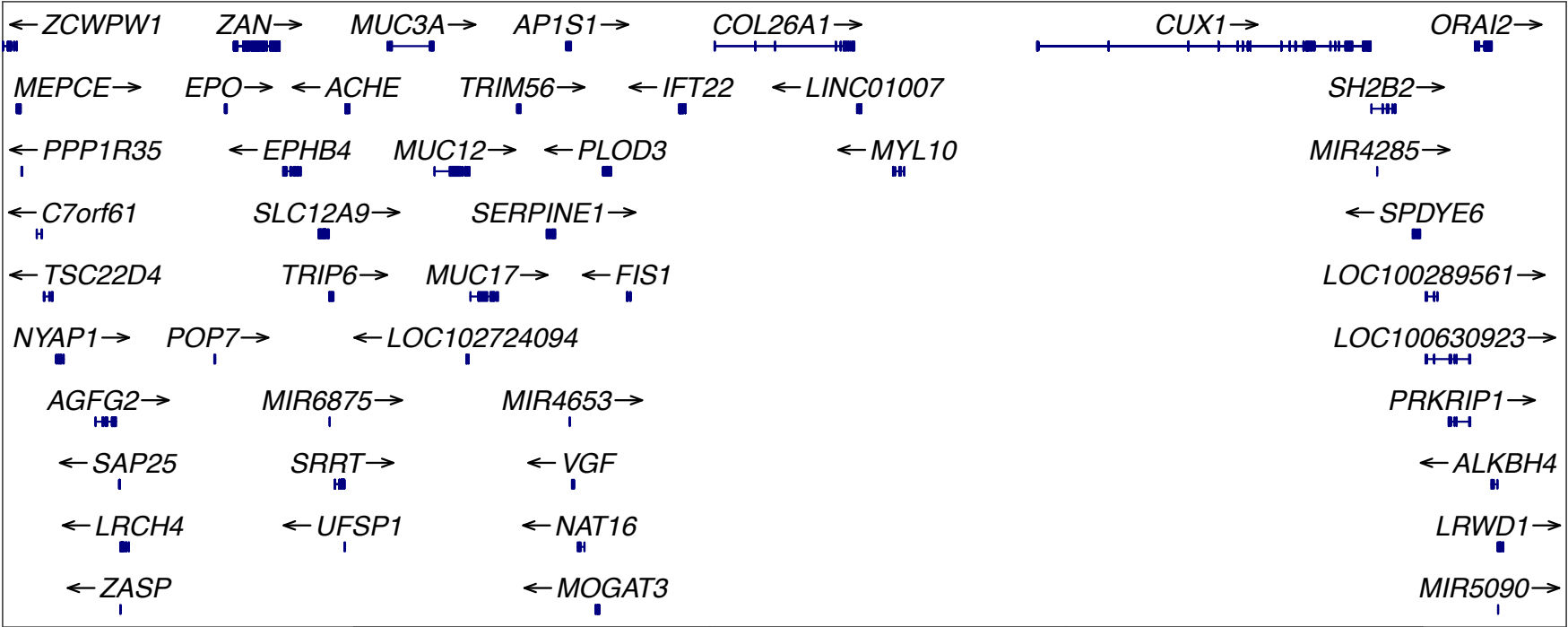
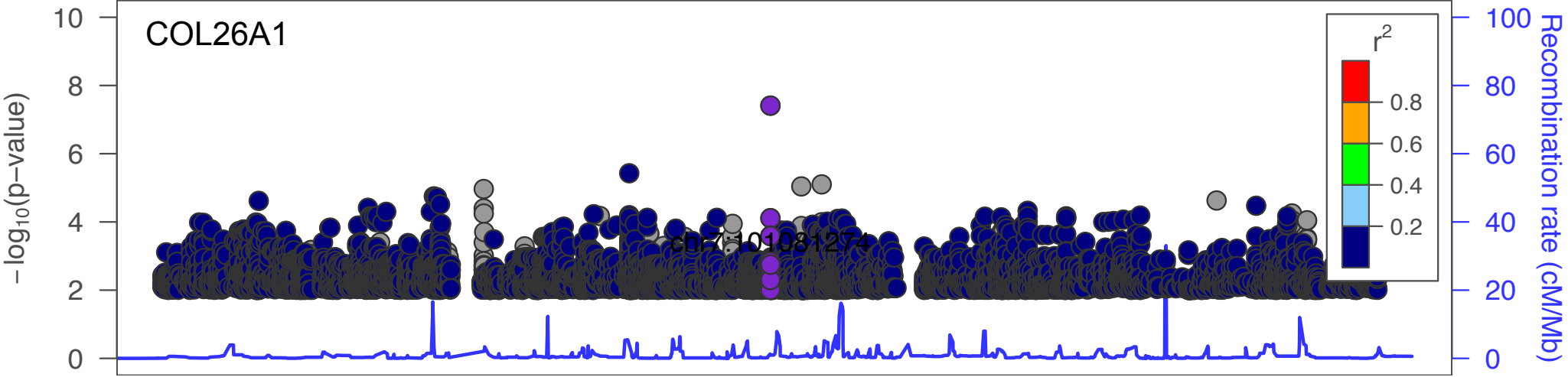
Plotted SNPs



Plotted SNPs



Plotted SNPs

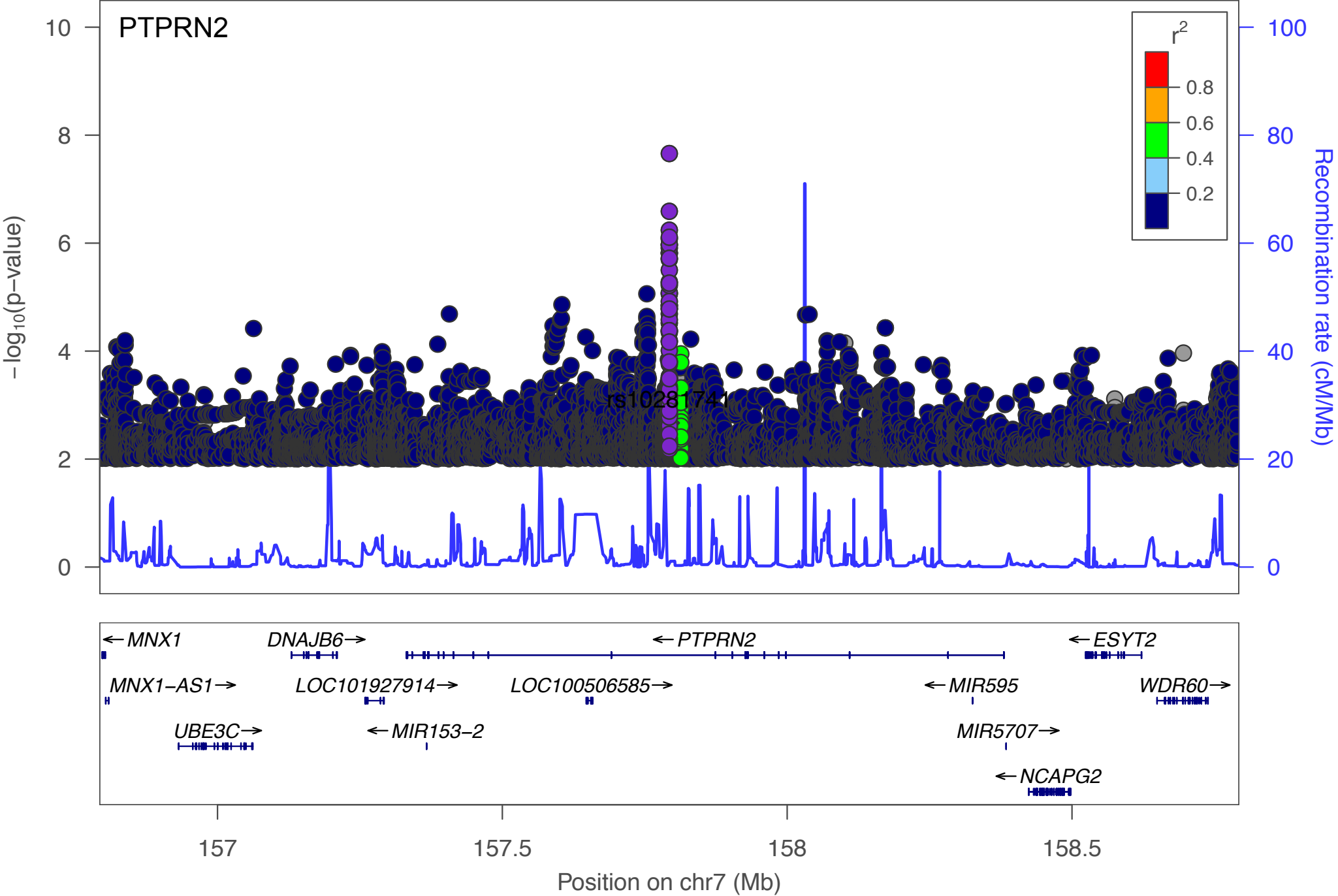


17 genes
omitted

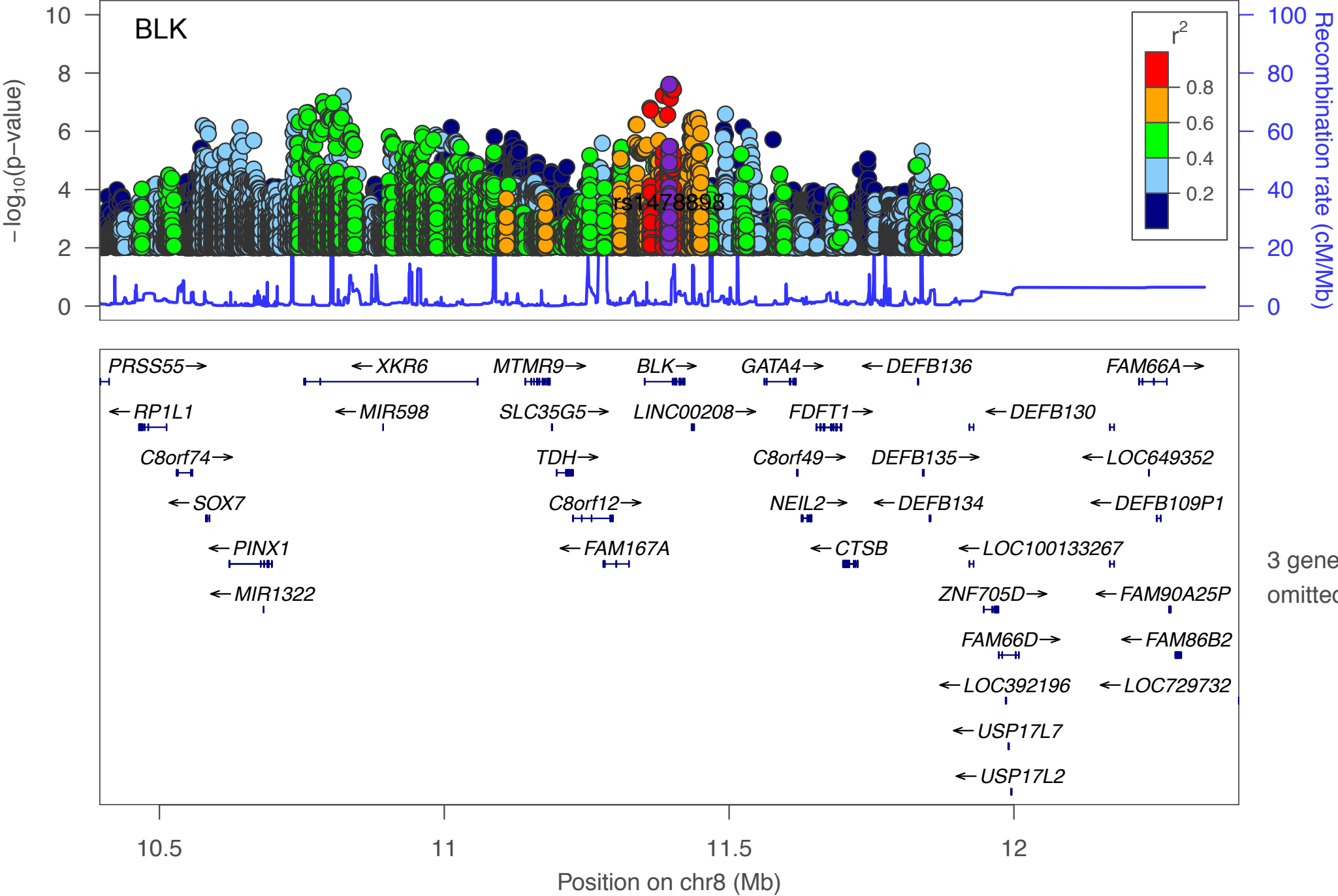
100.5 101 101.5 102

Position on chr7 (Mb)

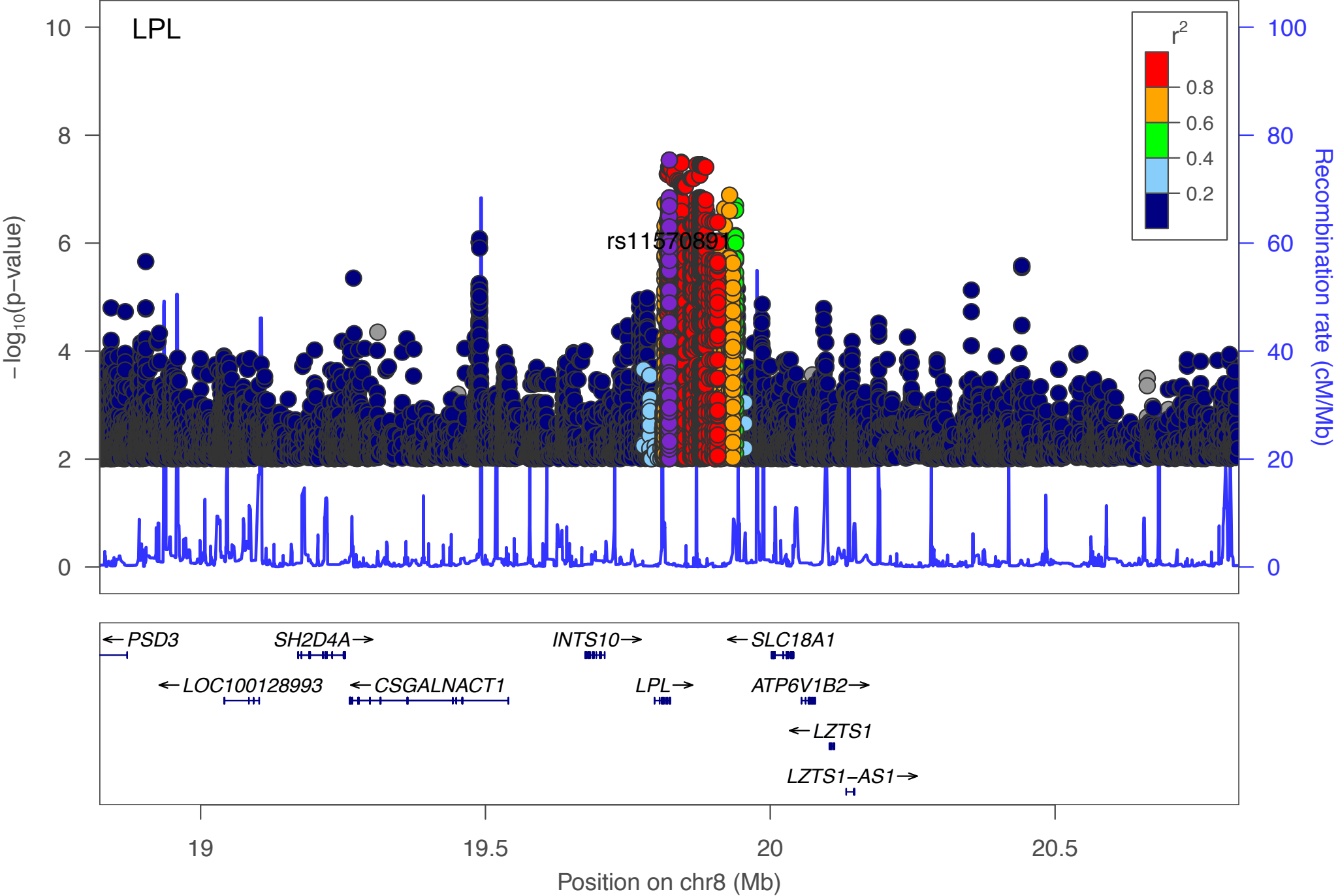
Plotted SNPs

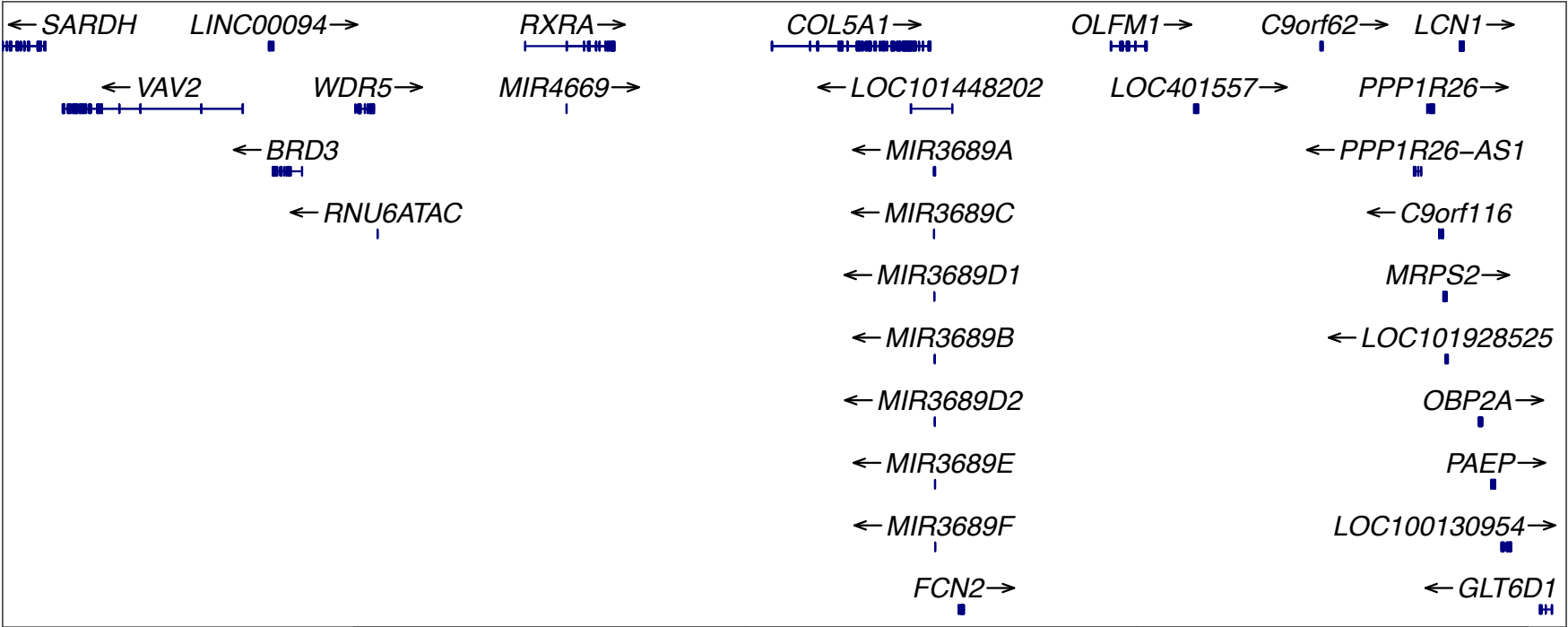
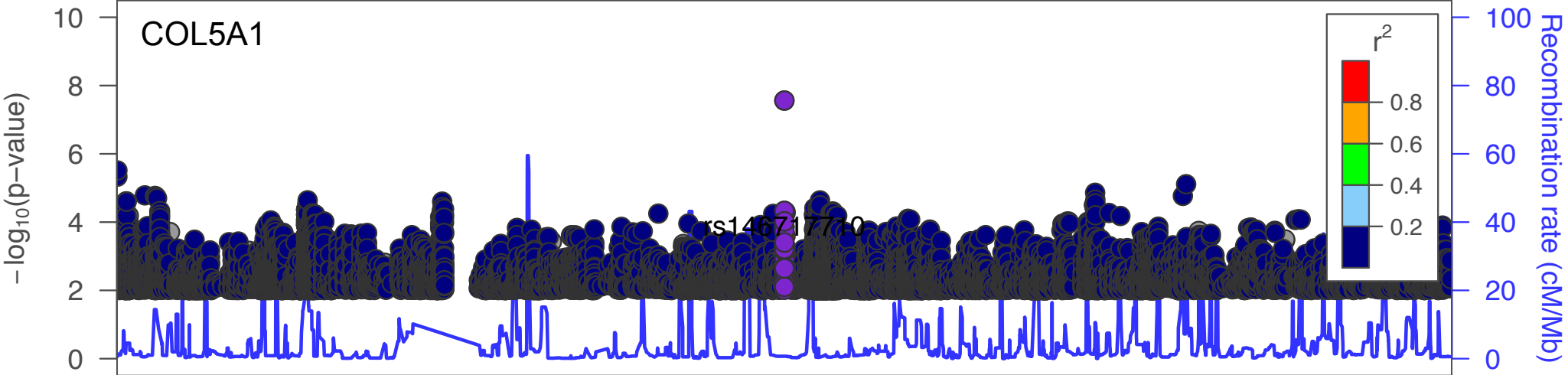


Plotted SNPs

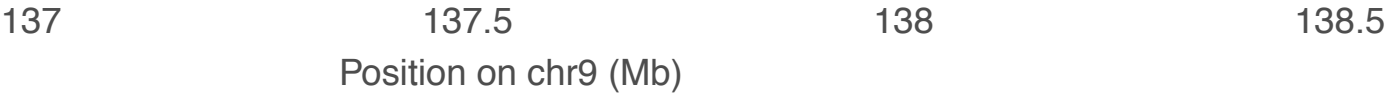


Plotted SNPs

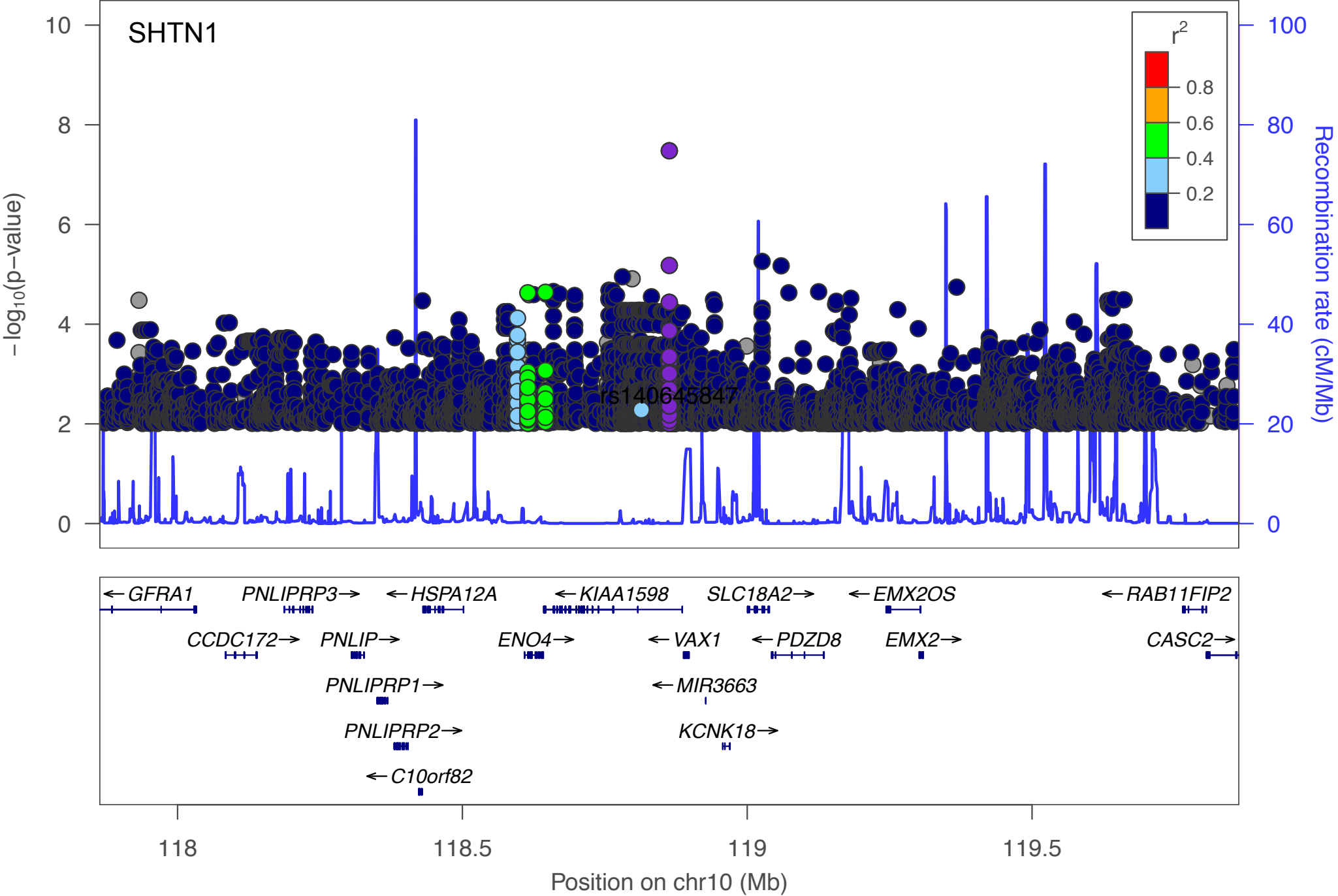


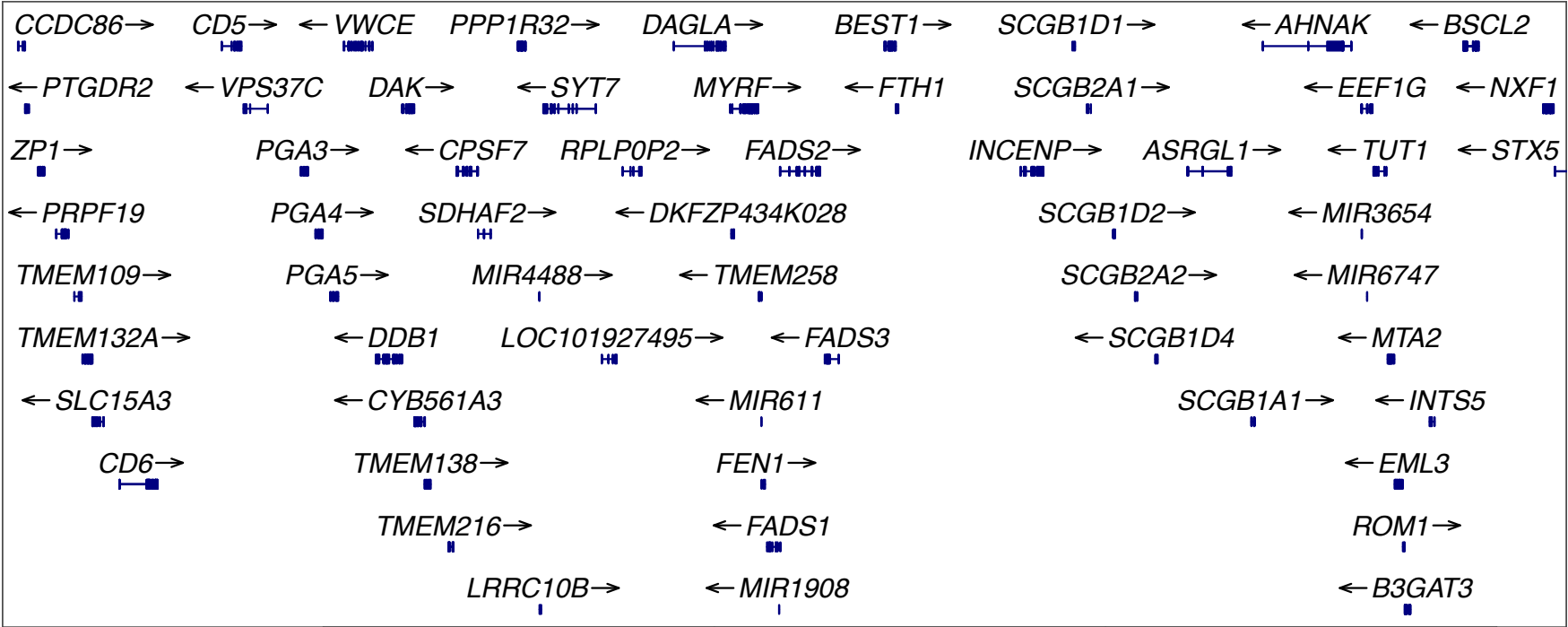
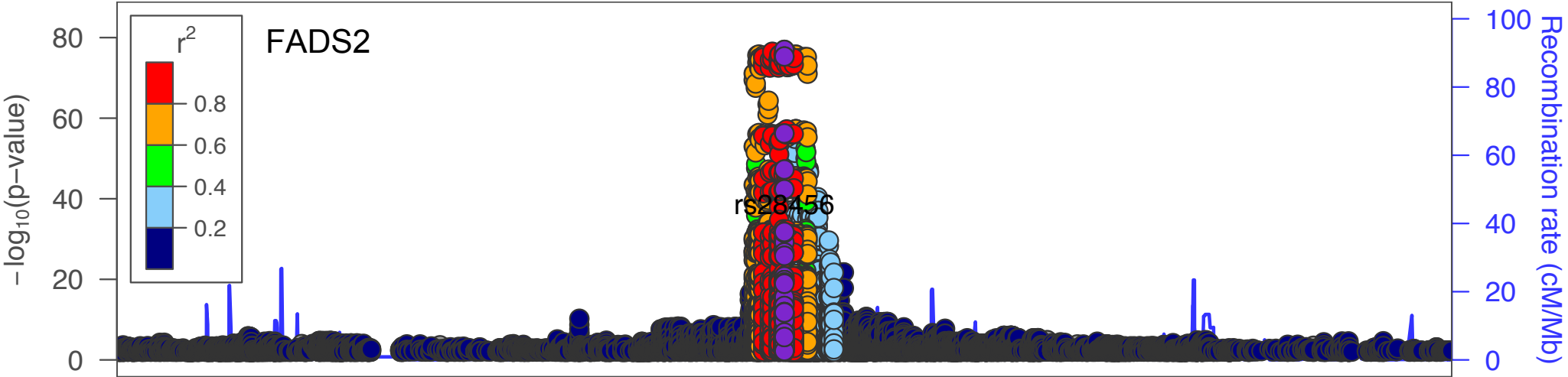


1 gene
omitted

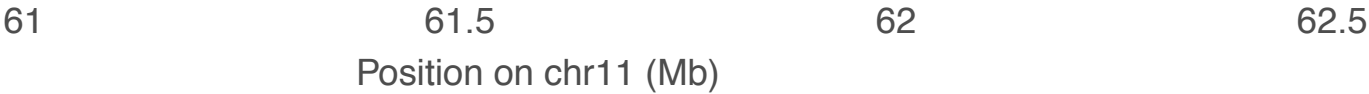


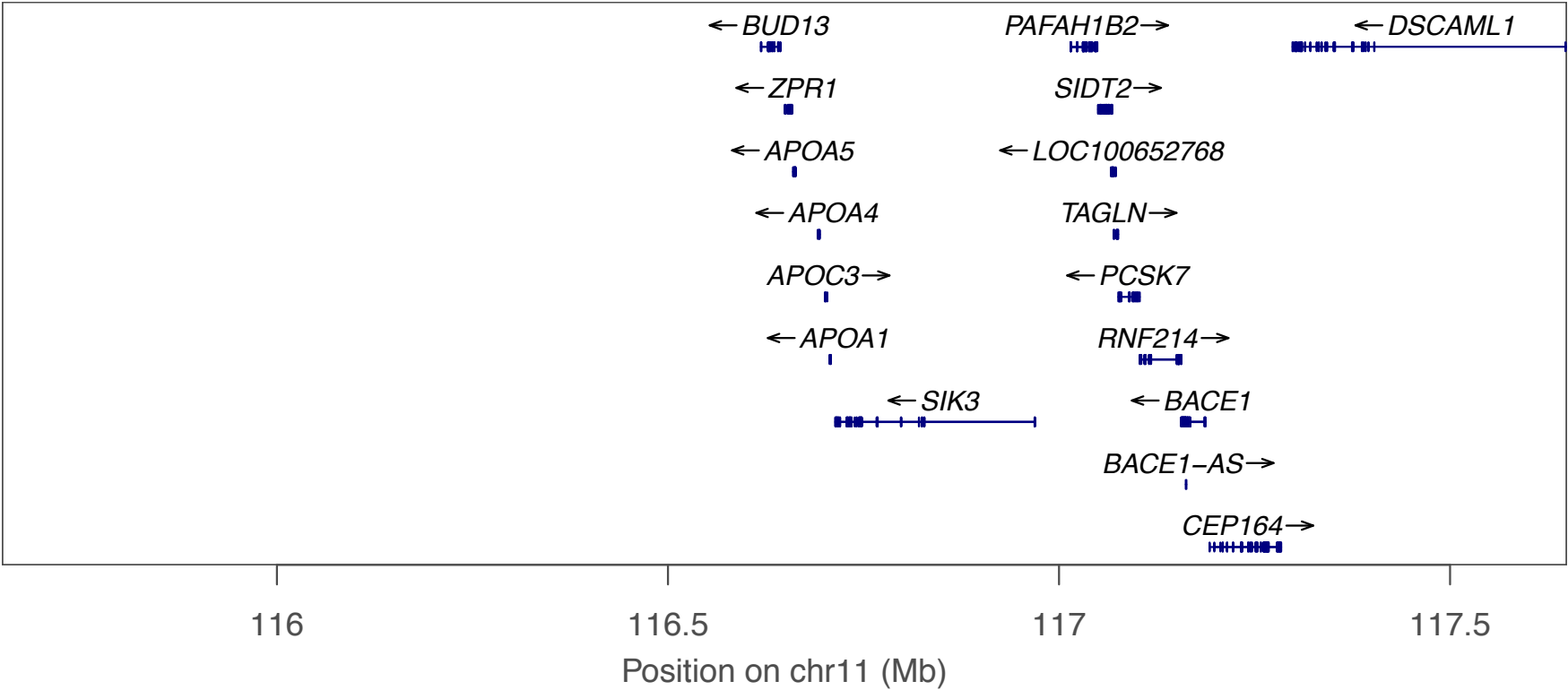
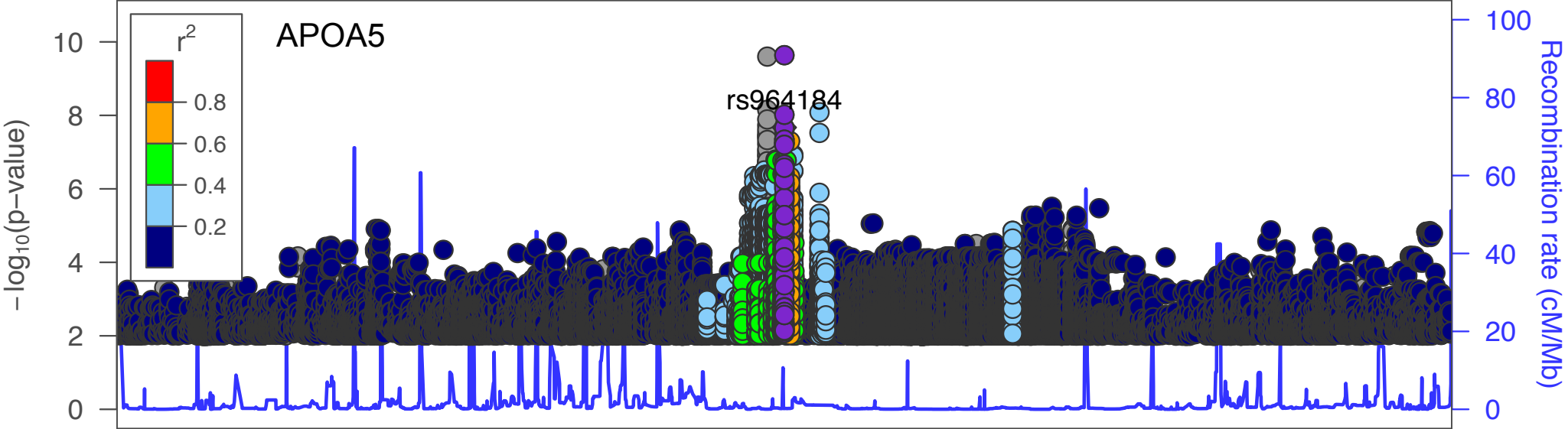
Plotted SNPs



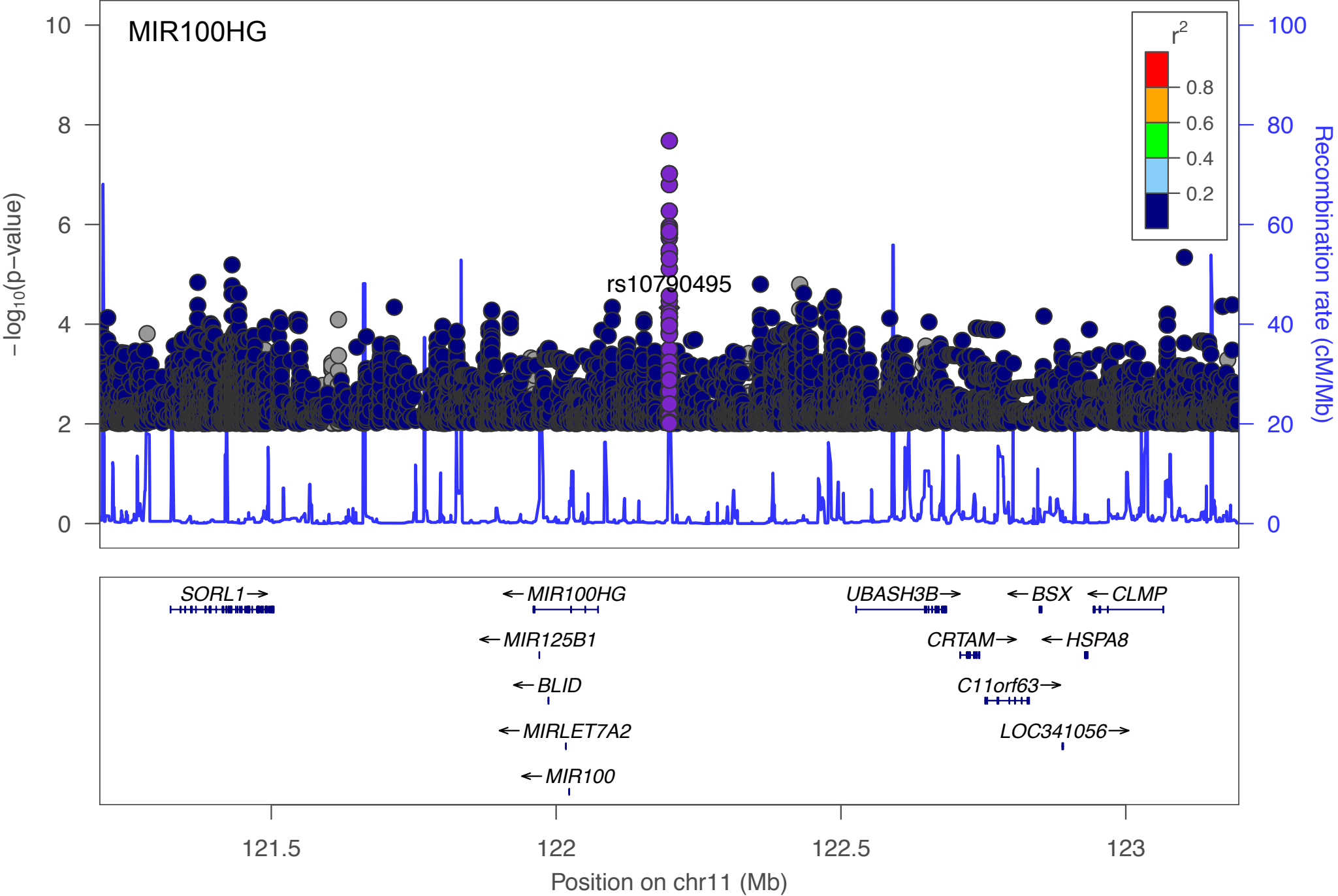


21 genes
omitted

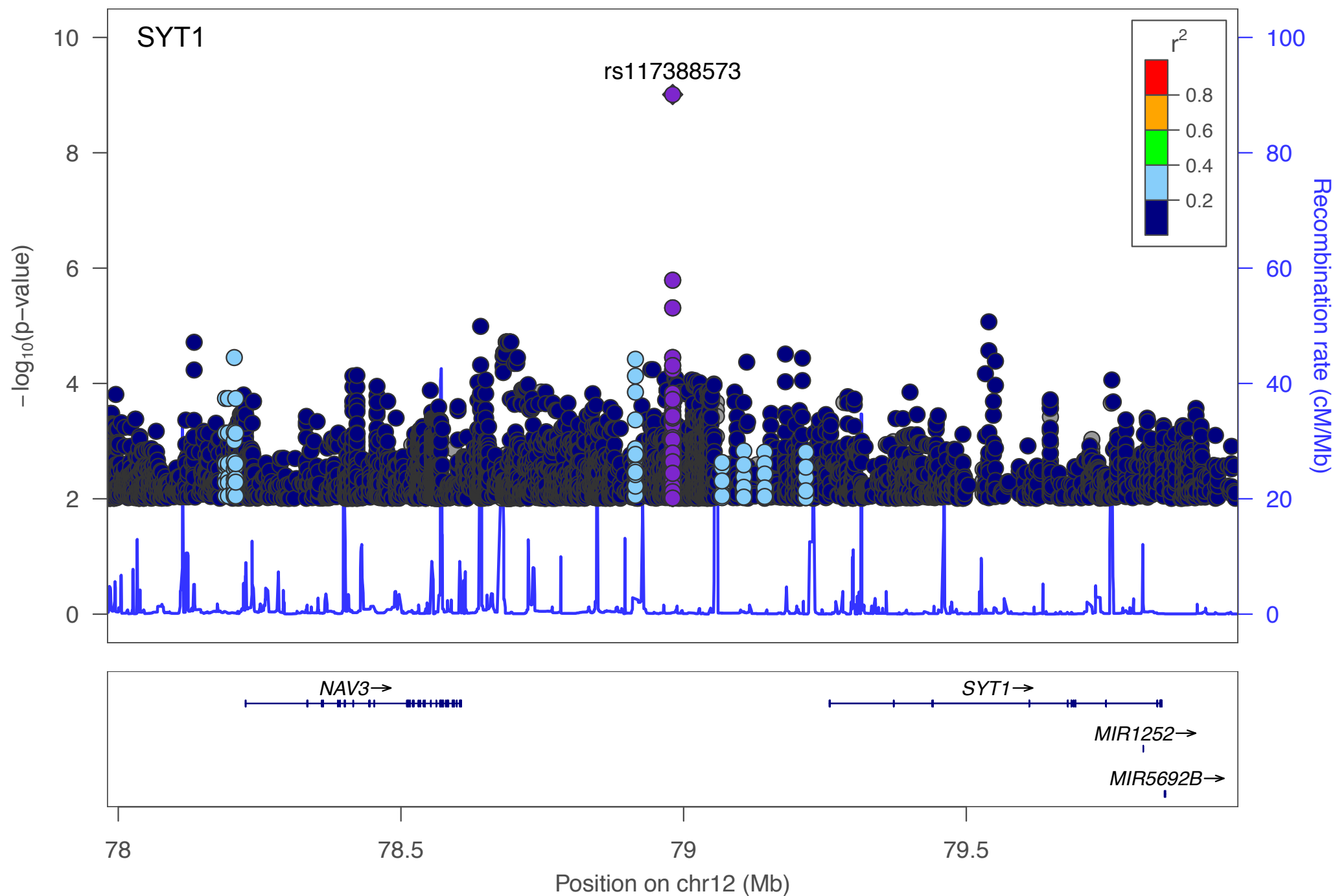




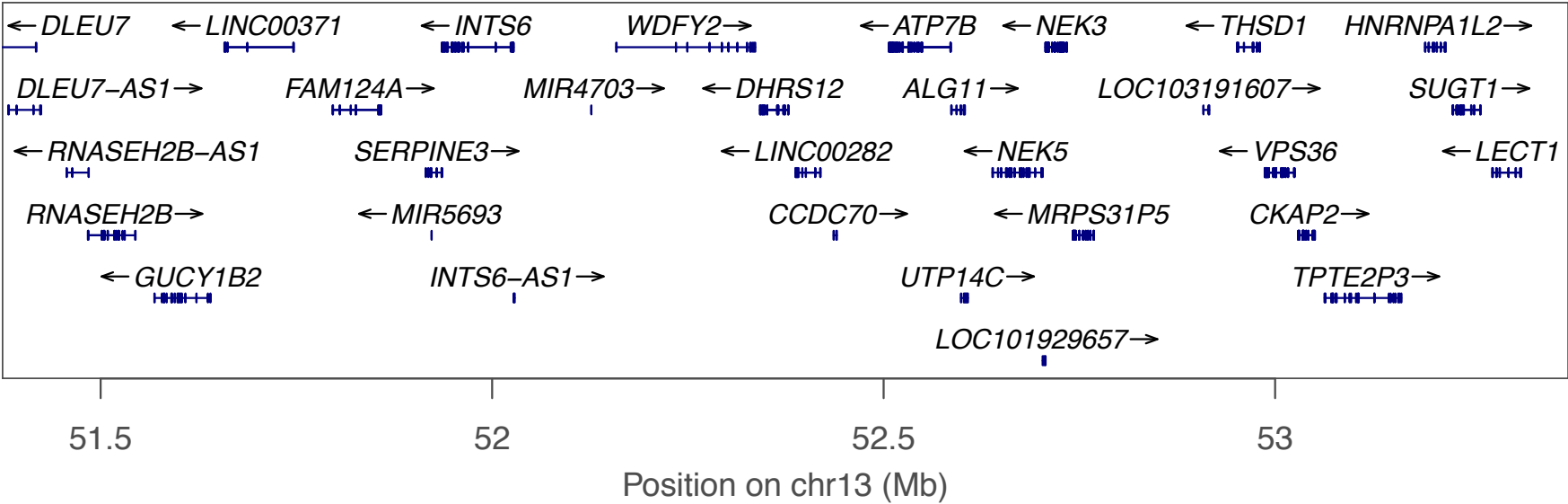
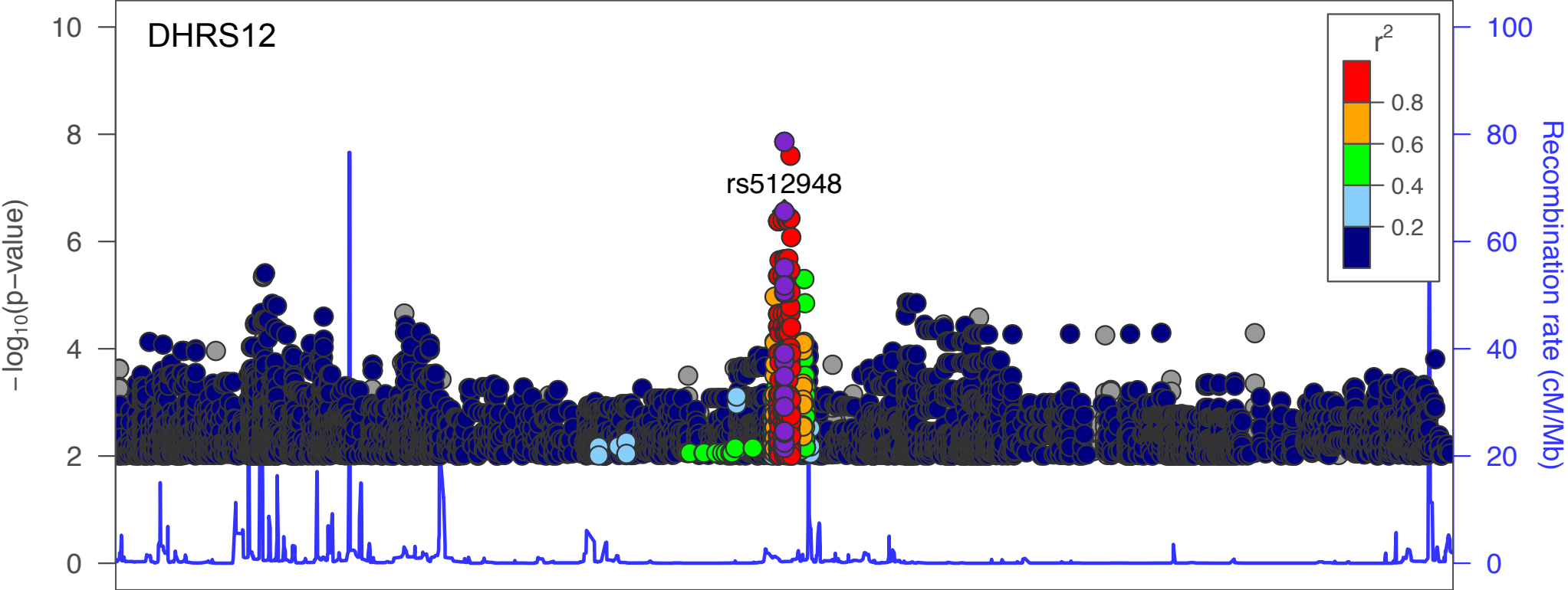
Plotted SNPs



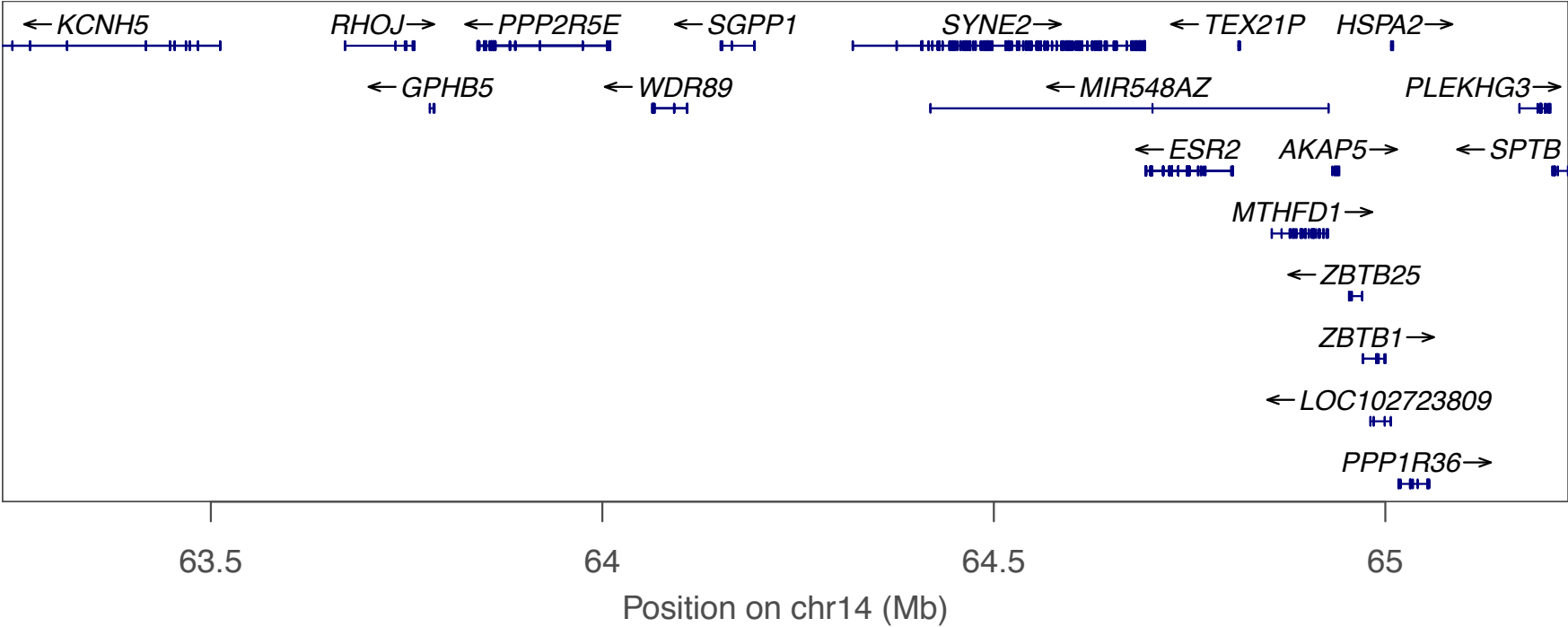
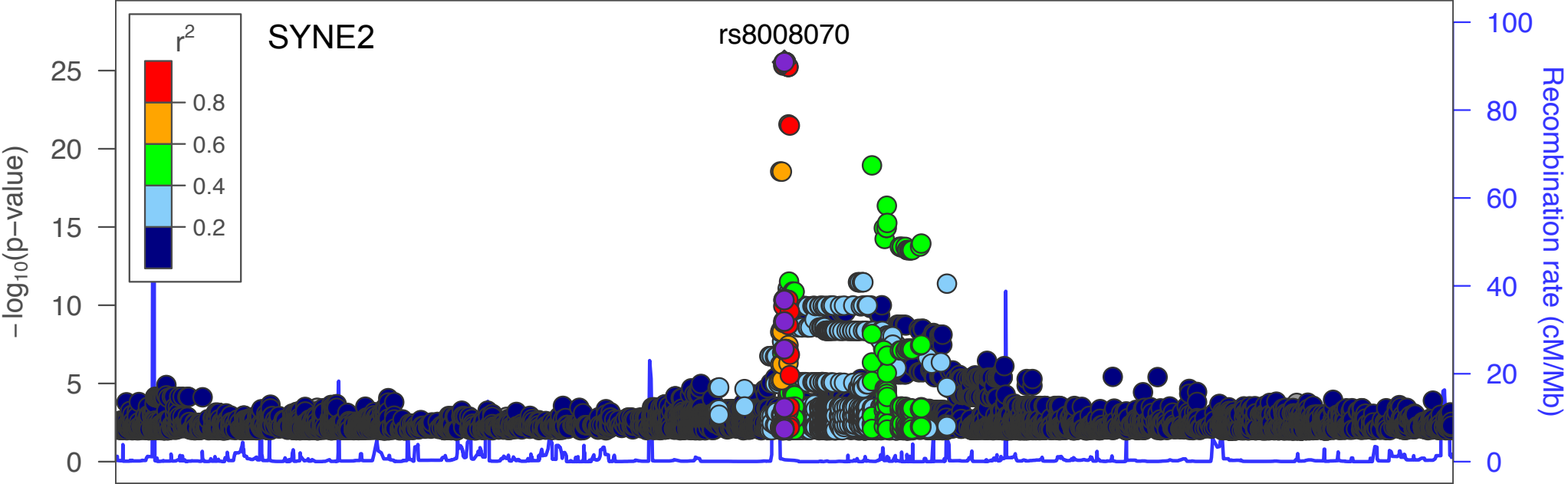
Plotted SNPs



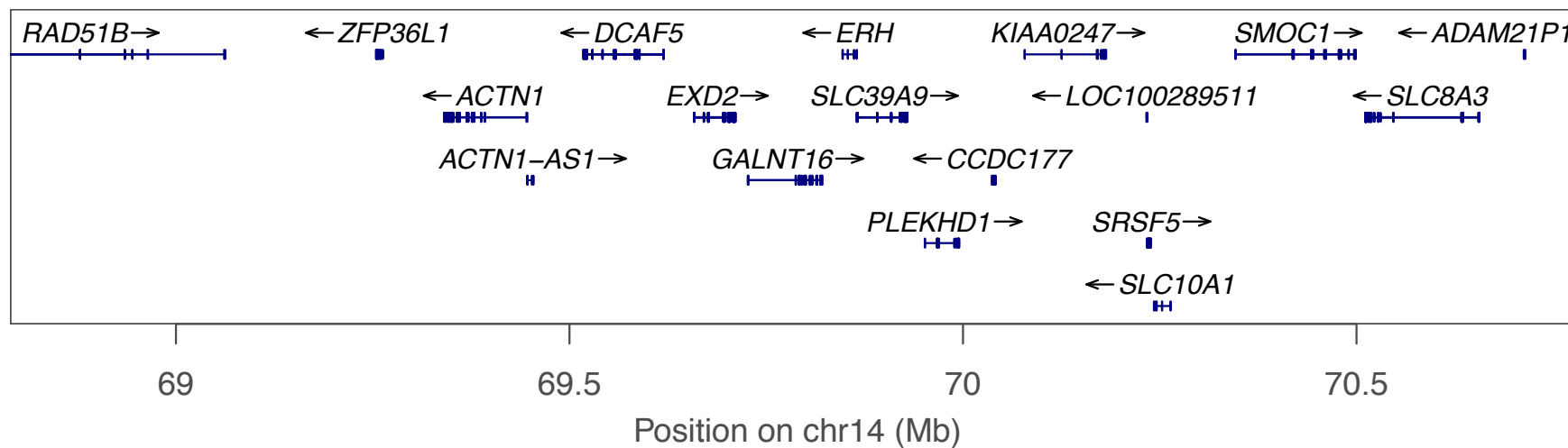
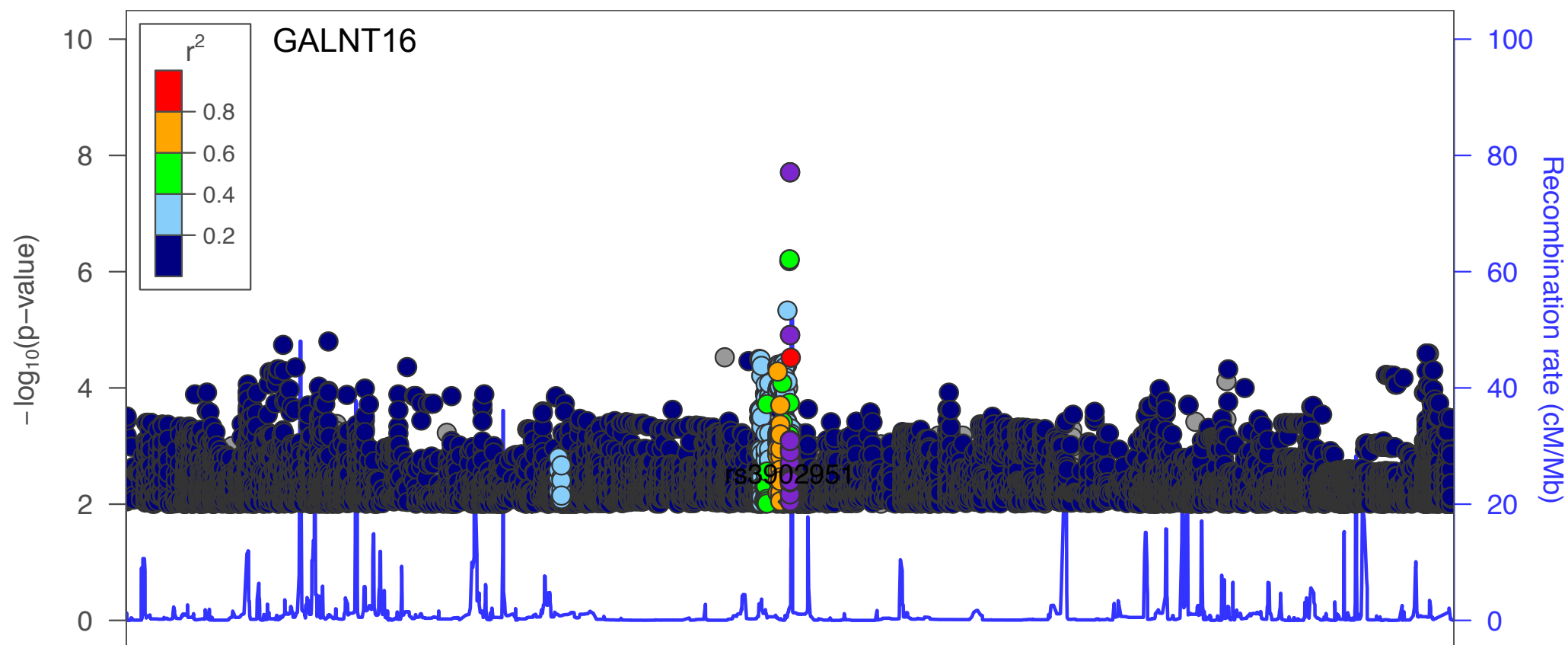
Plotted SNPs



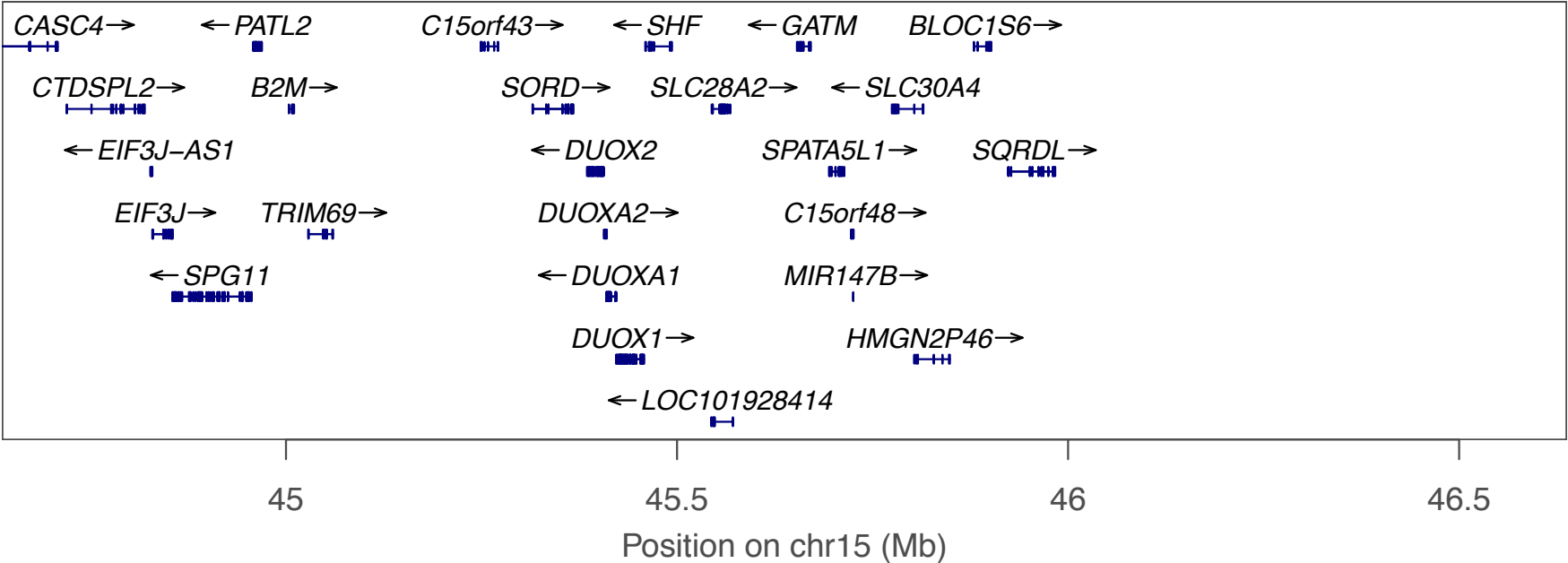
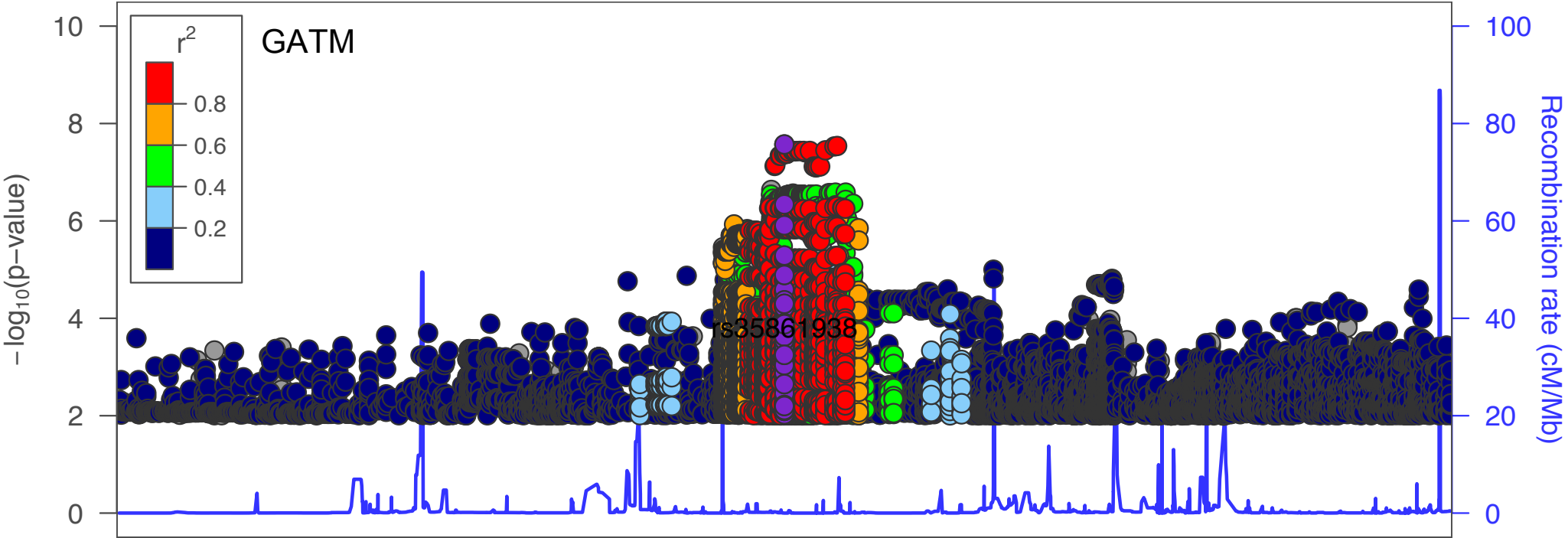
Plotted SNPs

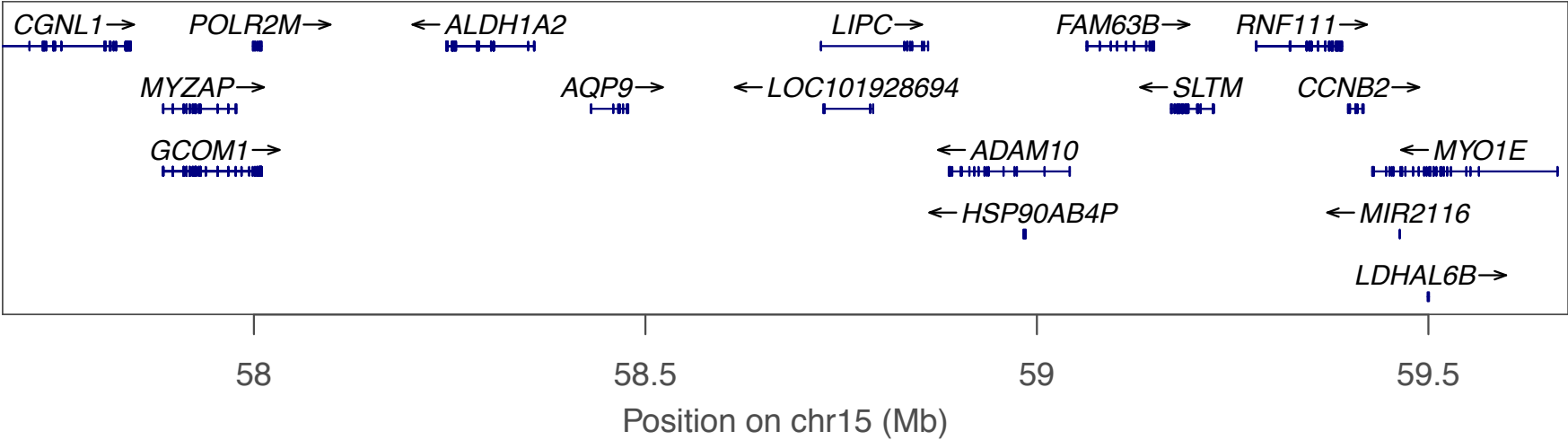
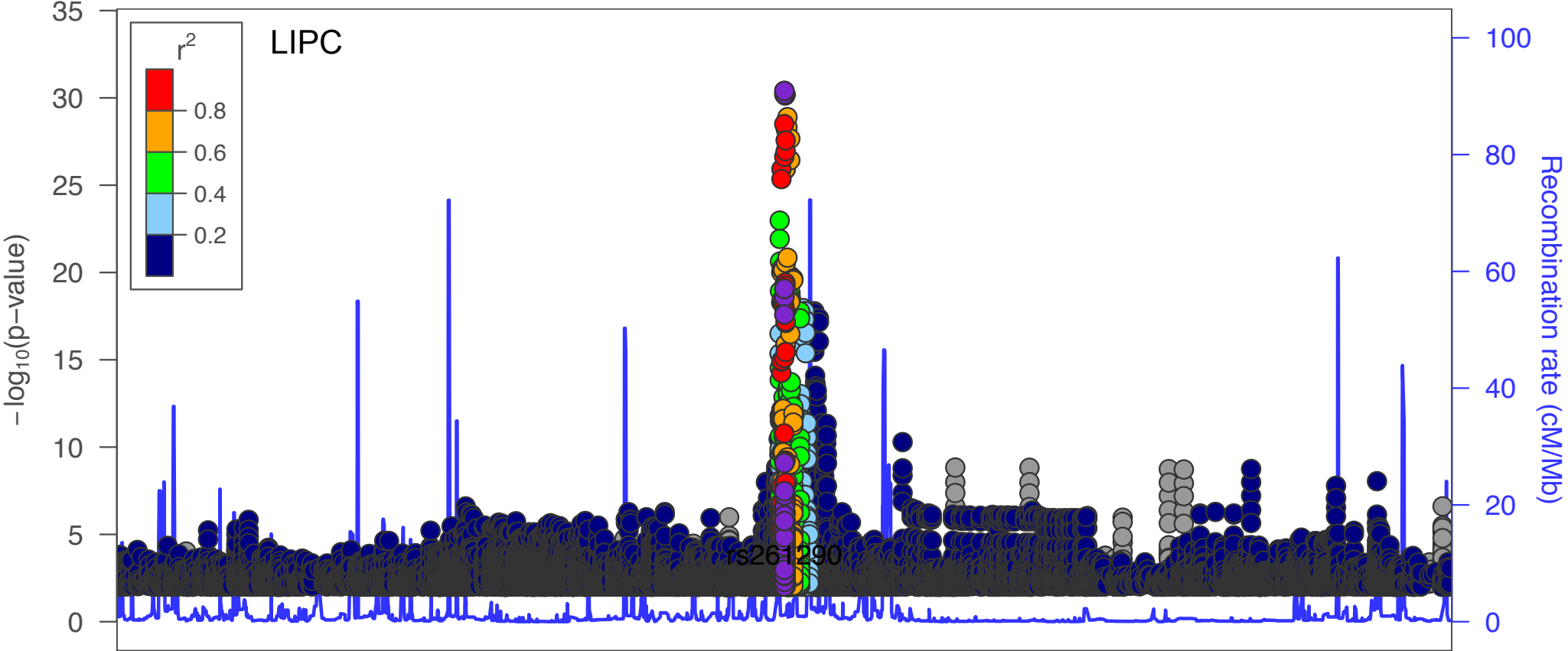


Plotted SNPs

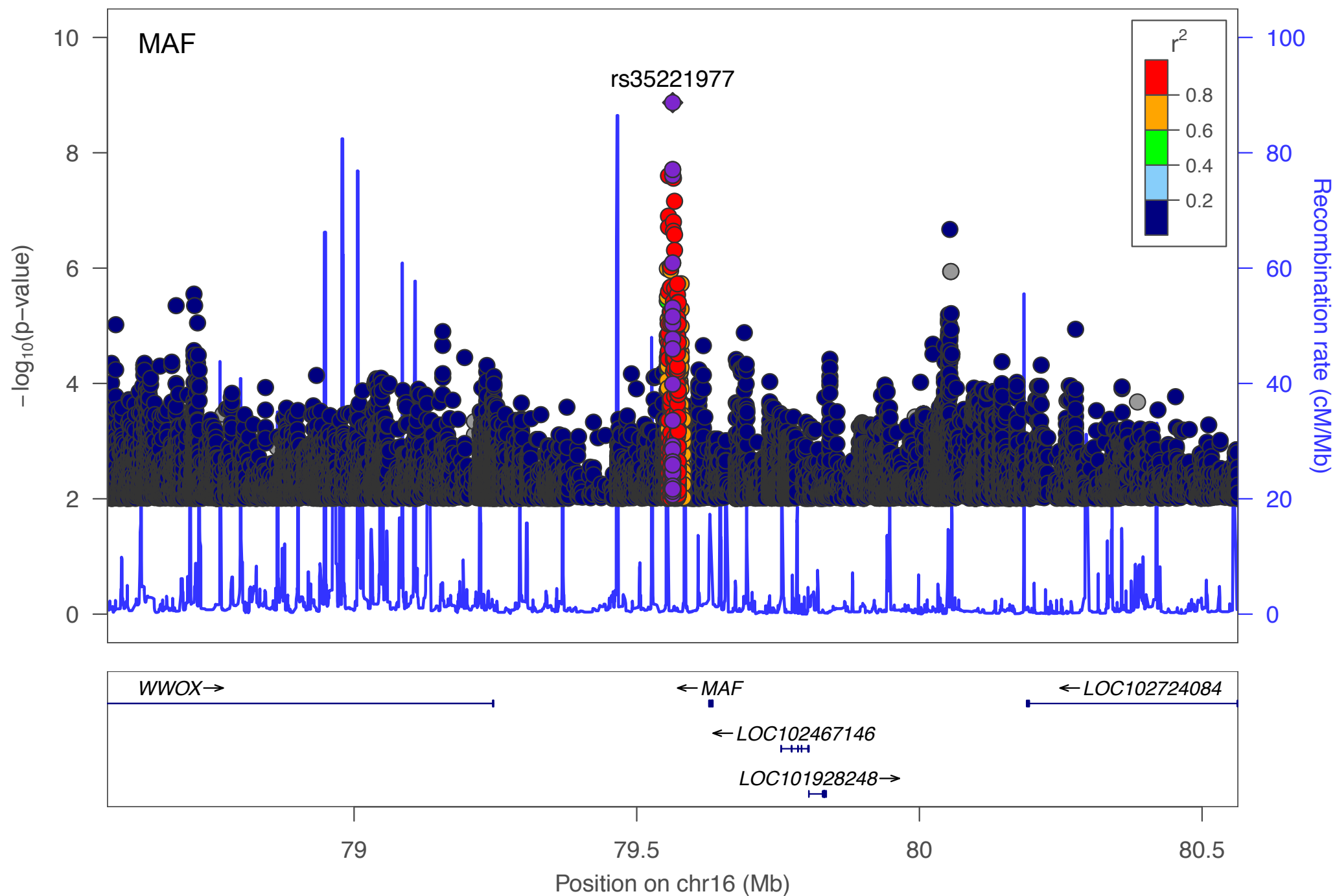


Plotted SNPs

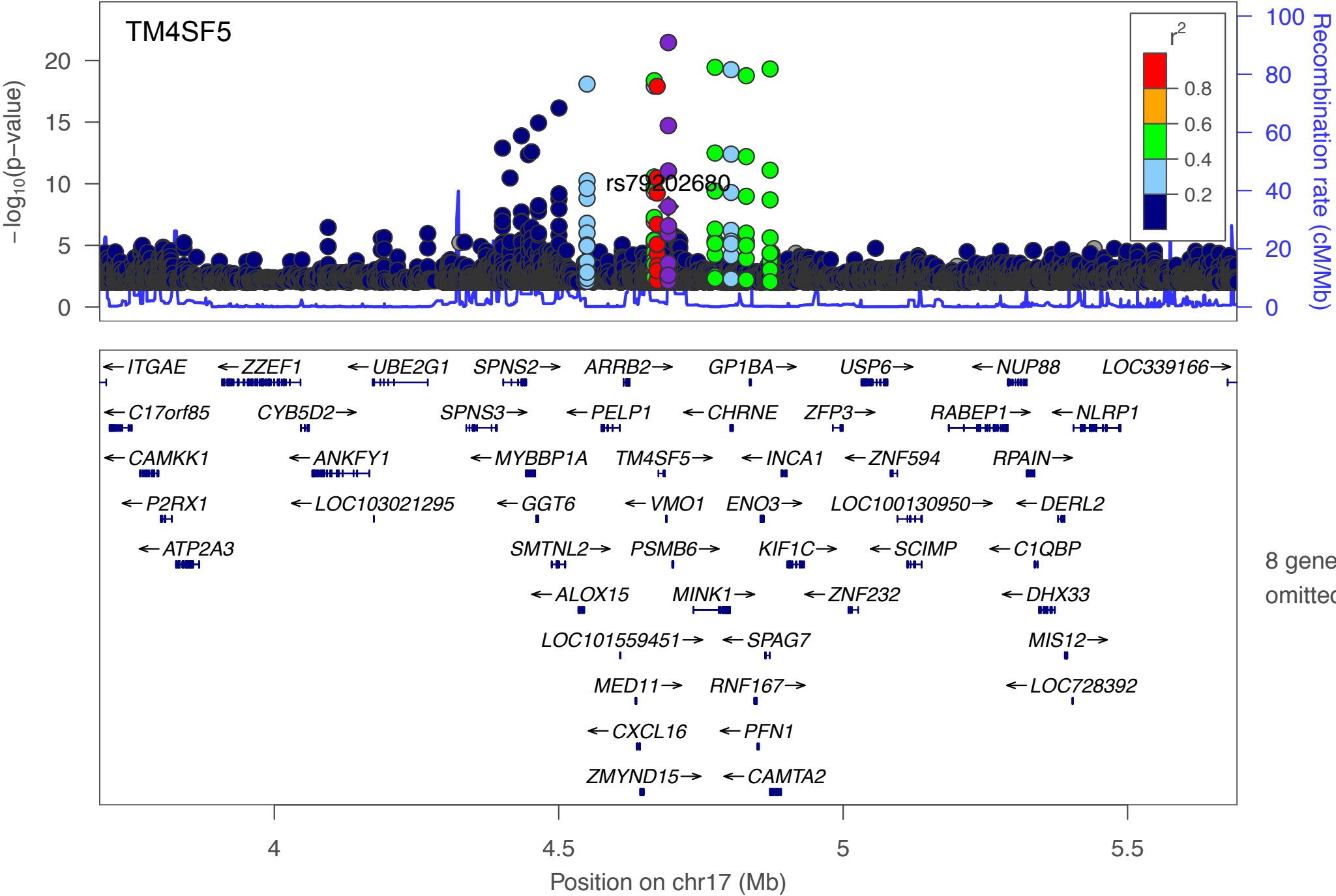


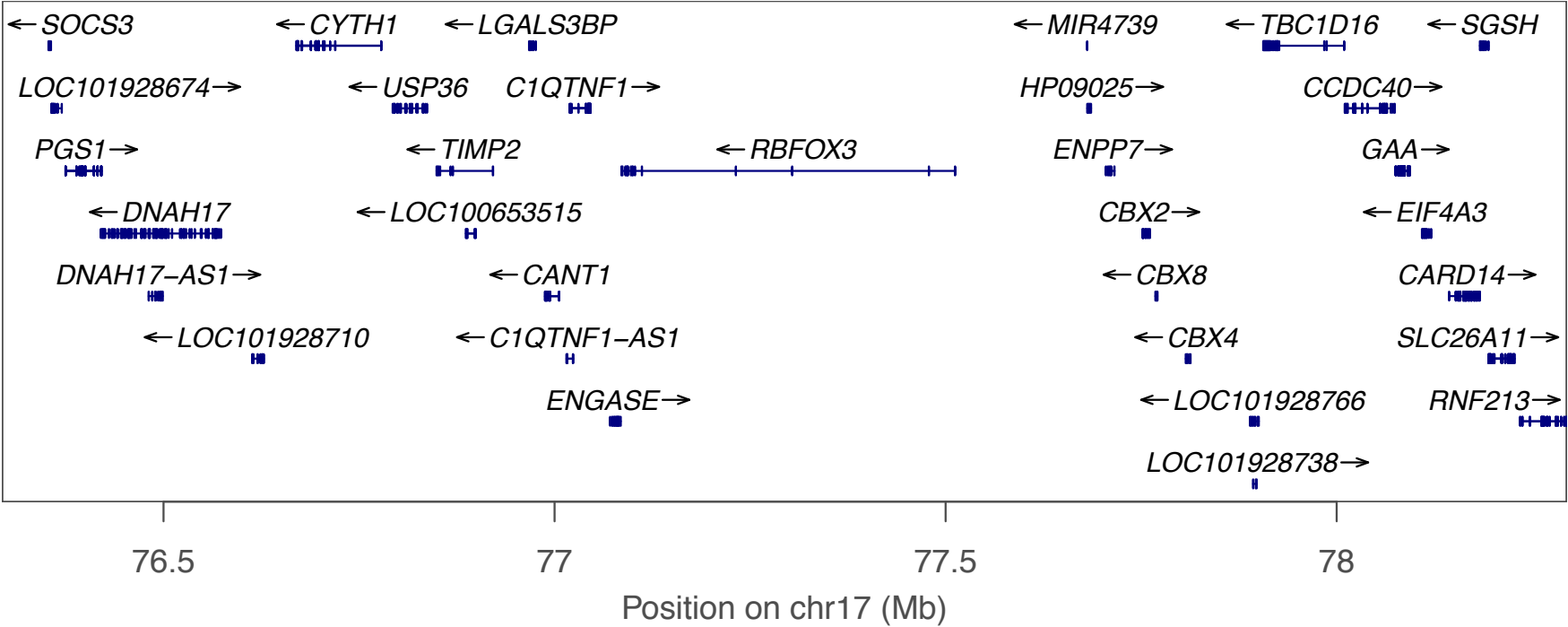
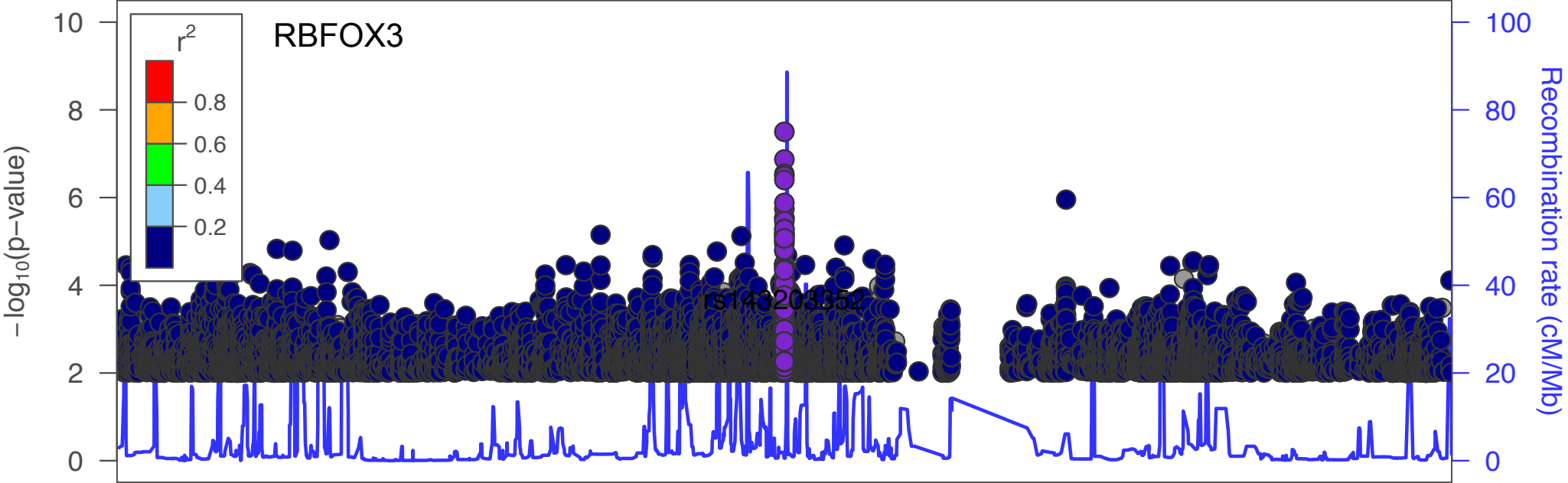


Plotted SNPs

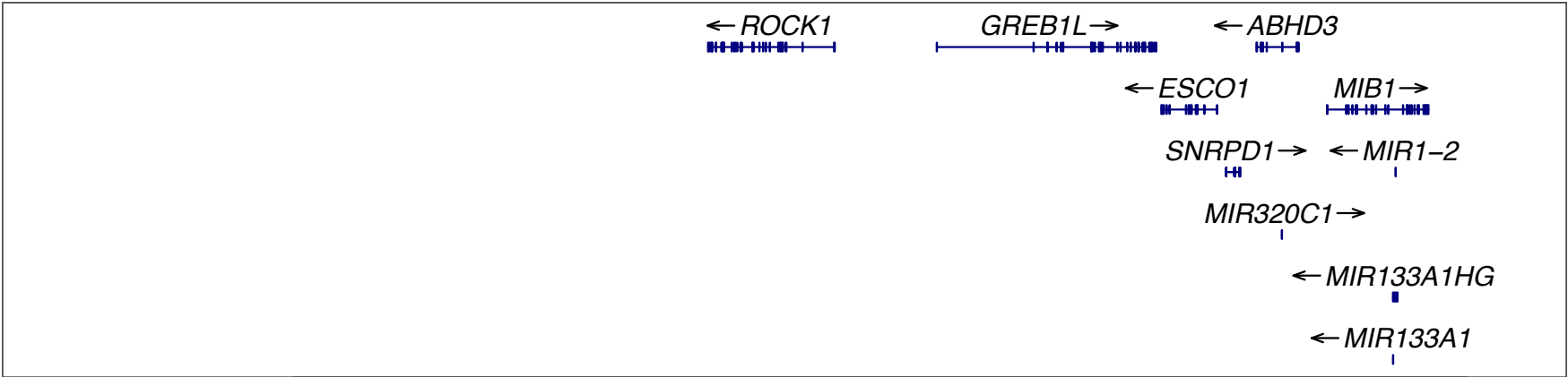
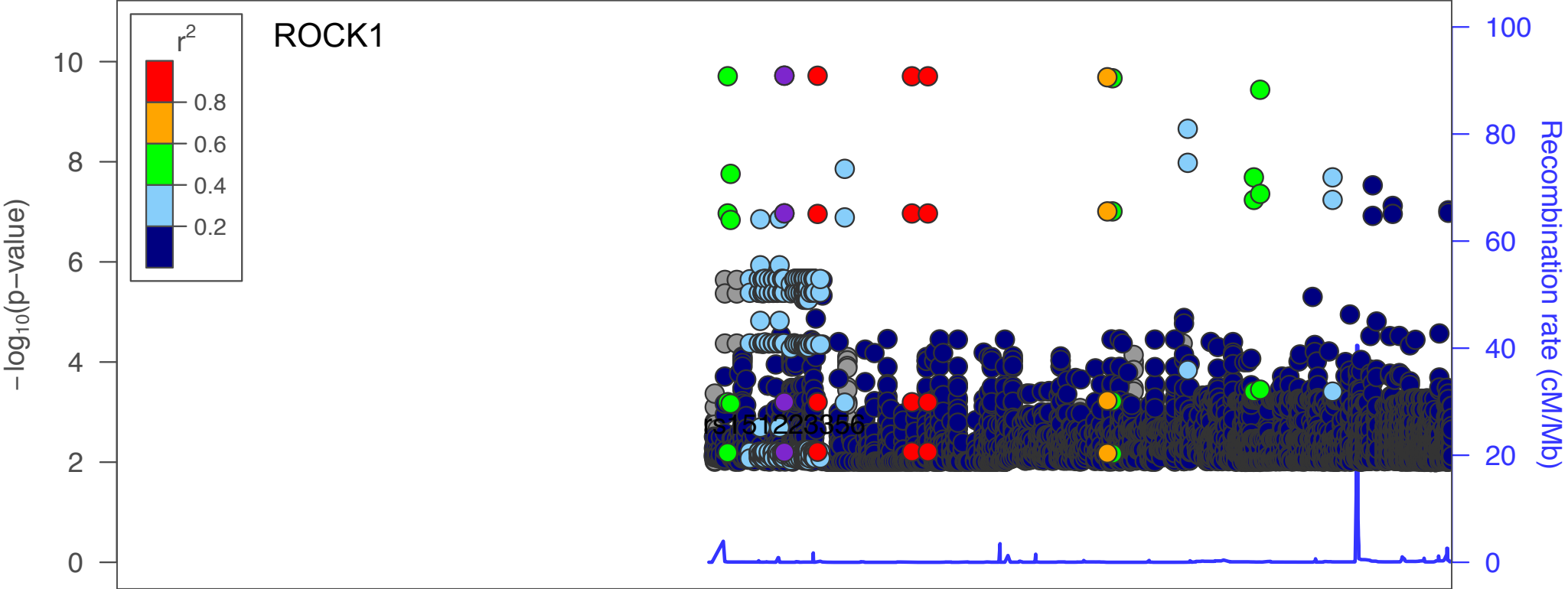


Plotted SNPs





Plotted SNPs



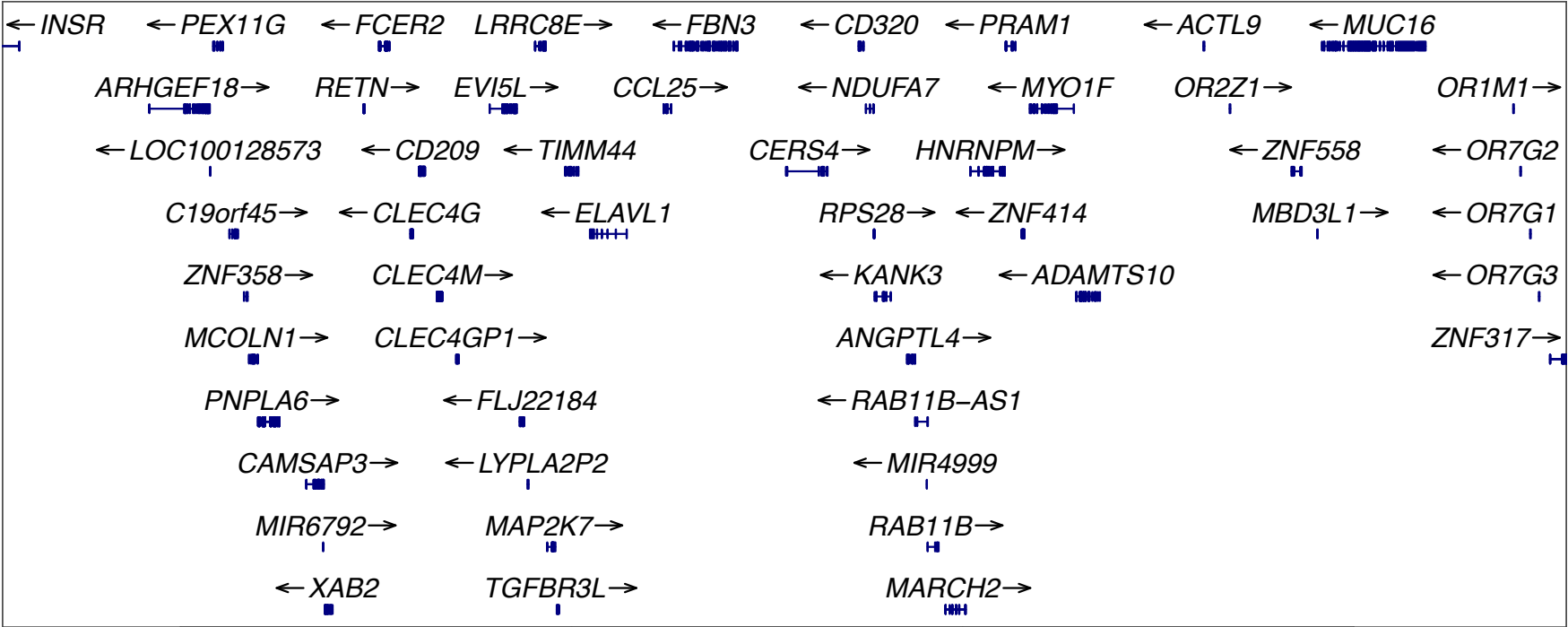
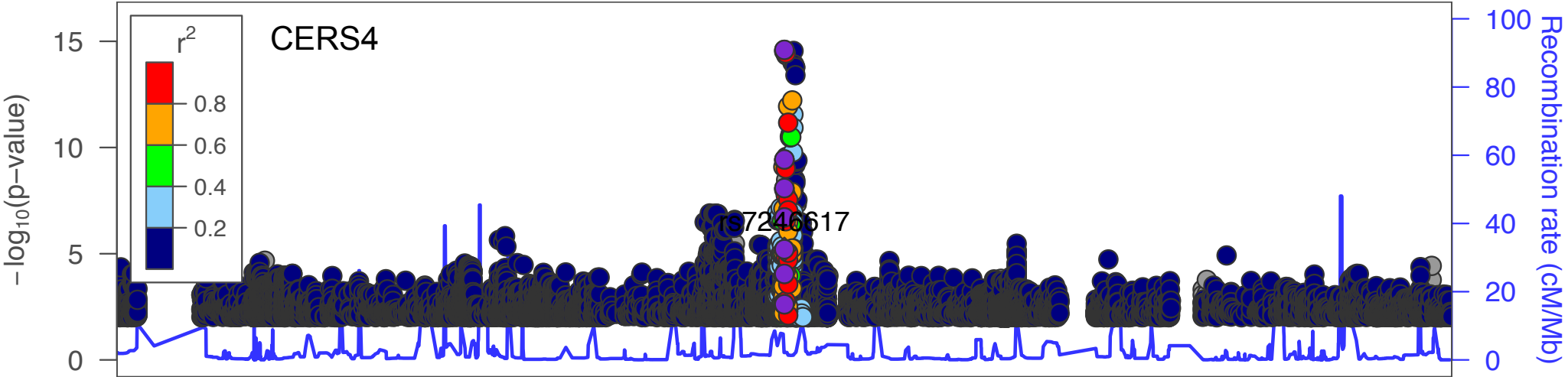
18

18.5

19

19.5

Position on chr18 (Mb)

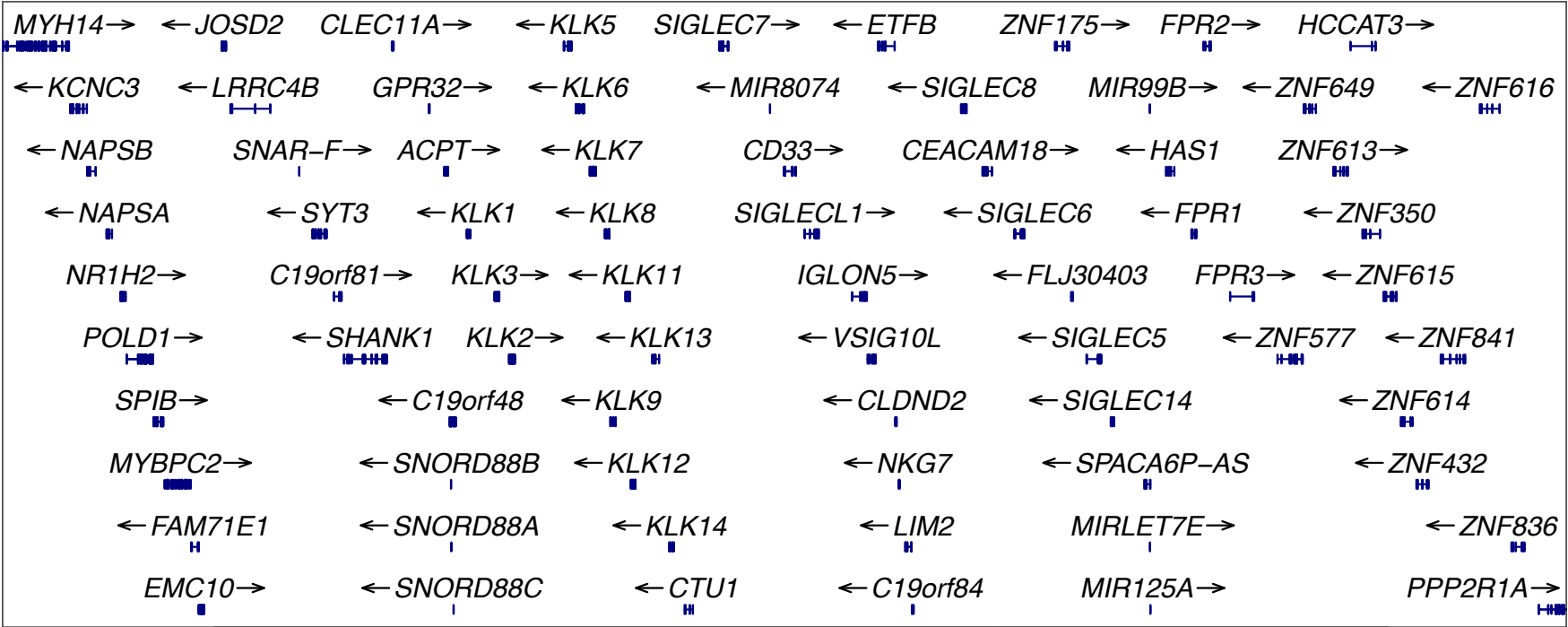
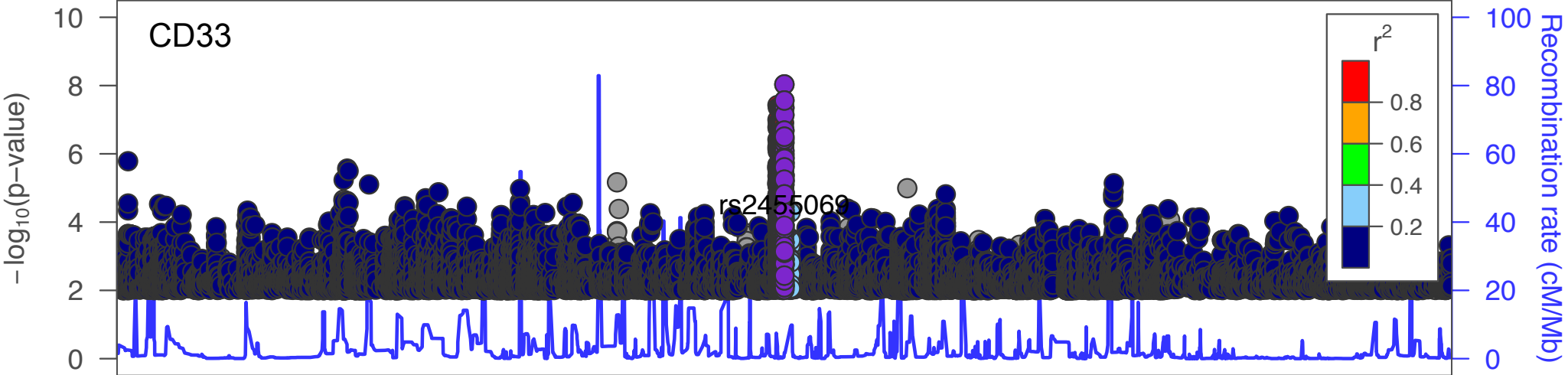


7 genes
omitted

7.5 8 8.5 9

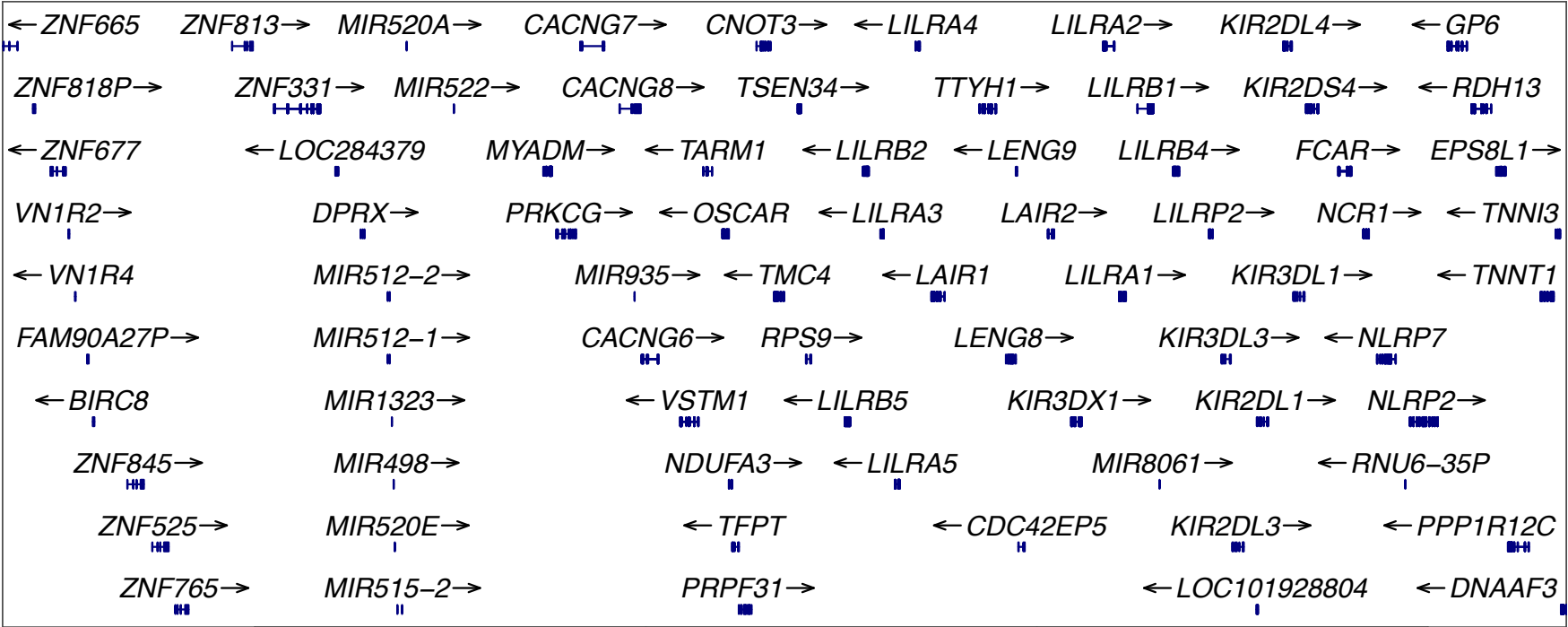
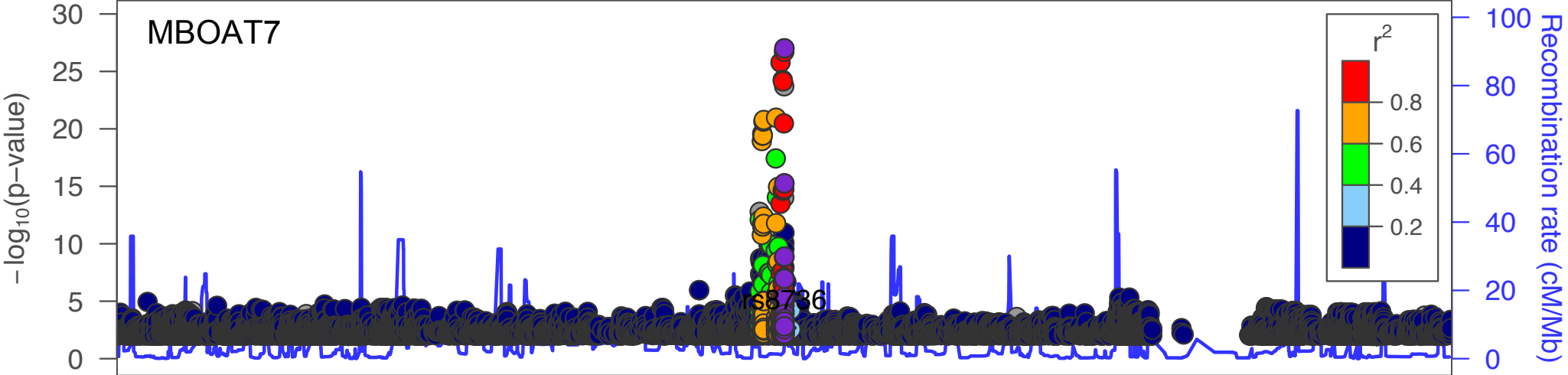
Position on chr19 (Mb)

Plotted SNPs



15 genes
omitted

51 51.5 52 52.5
Position on chr19 (Mb)



52 genes
omitted

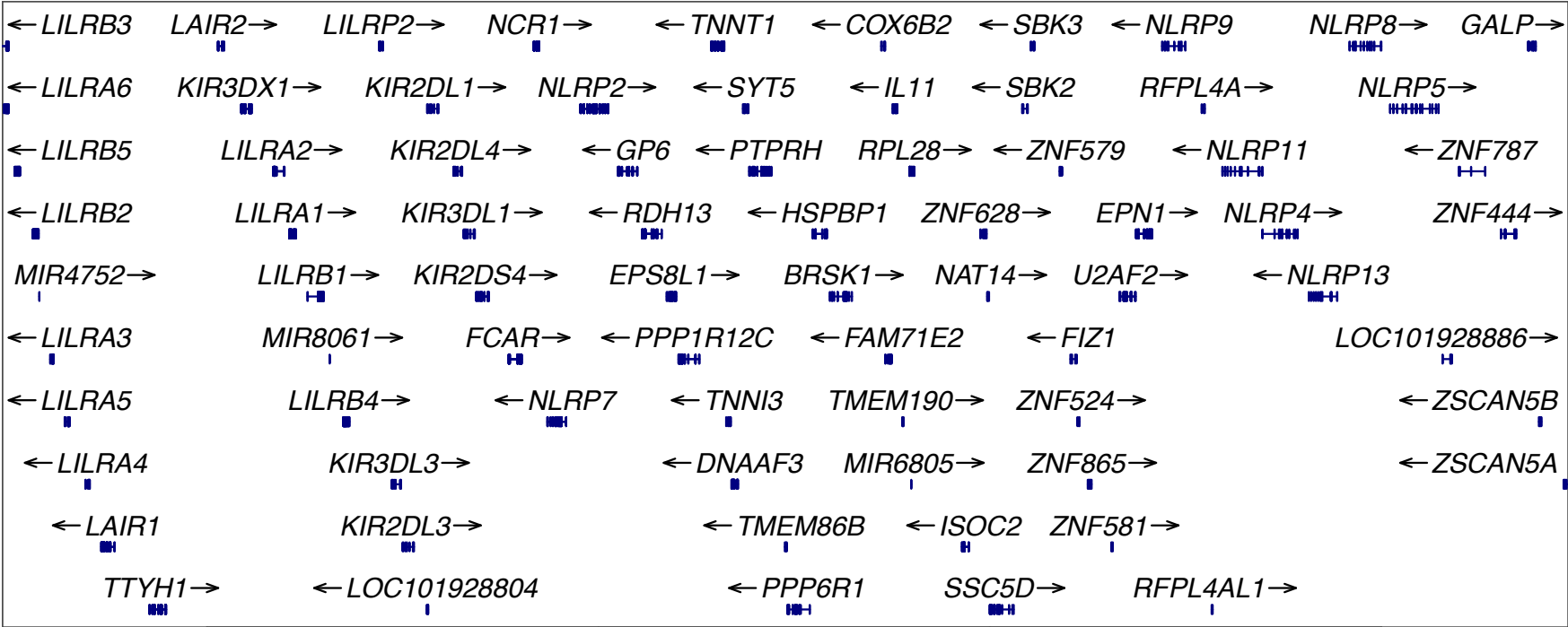
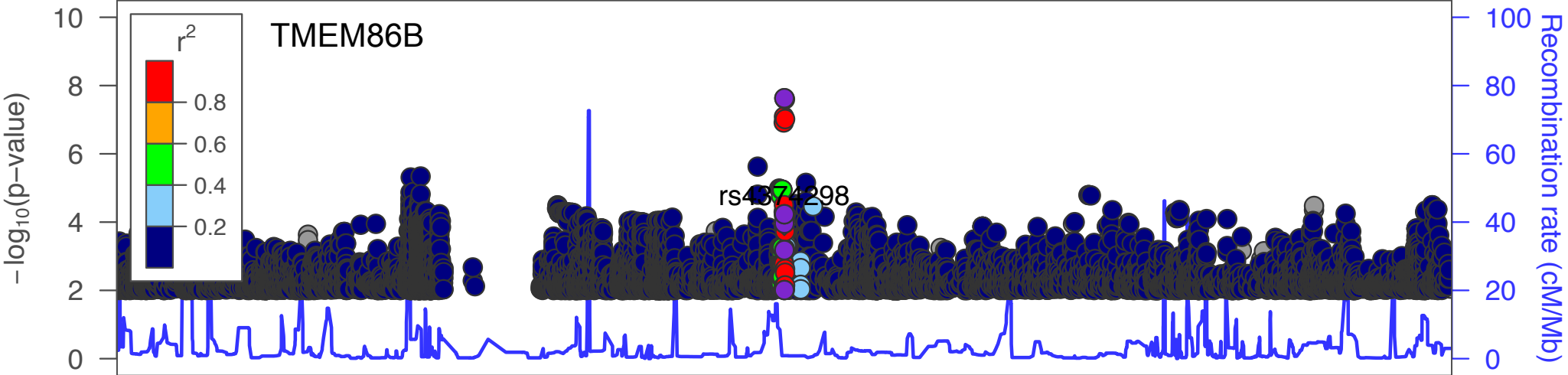
54

54.5

55

55.5

Position on chr19 (Mb)



17 genes
omitted

55

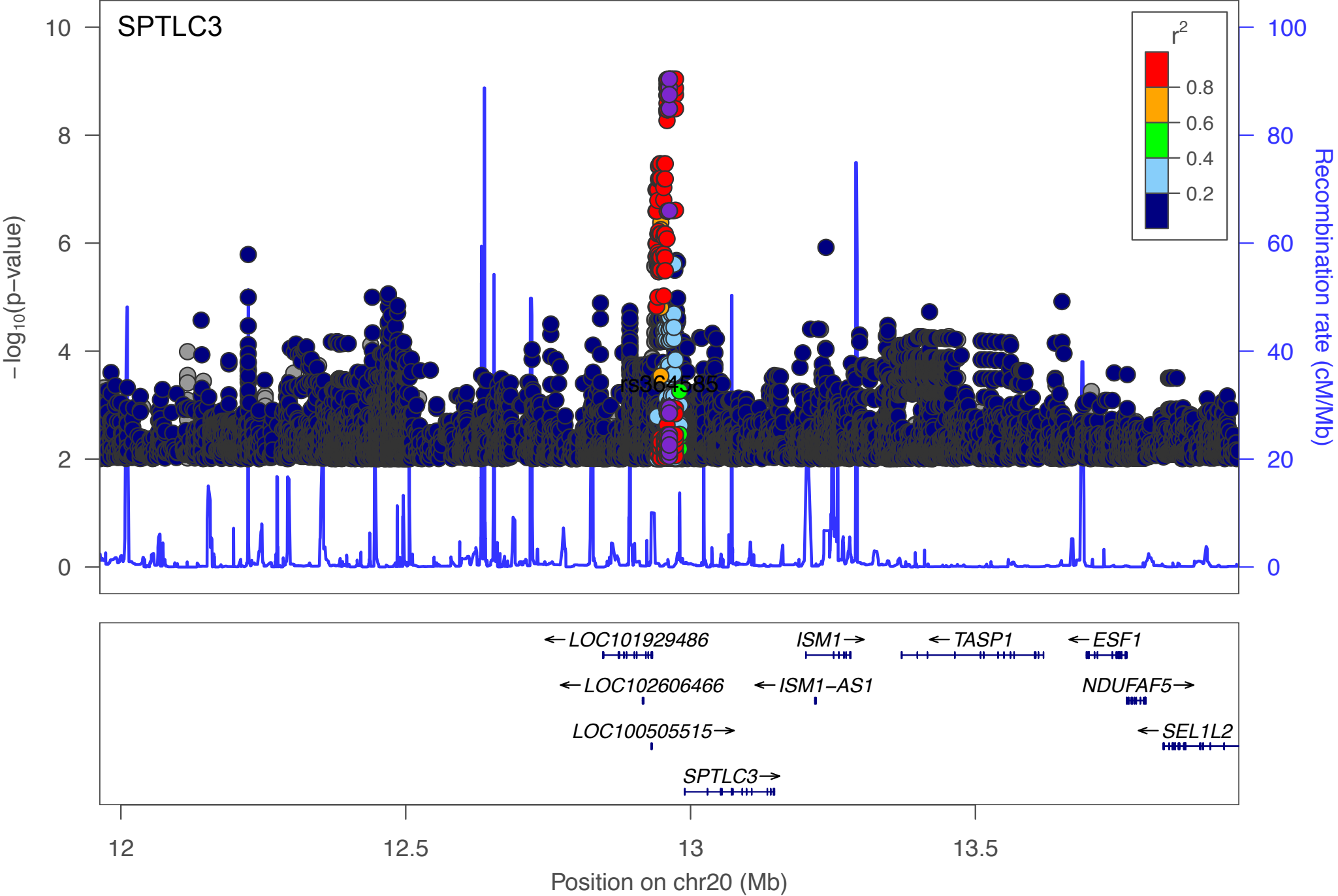
55.5

56

56.5

Position on chr19 (Mb)

Plotted SNPs



Plotted SNPs

