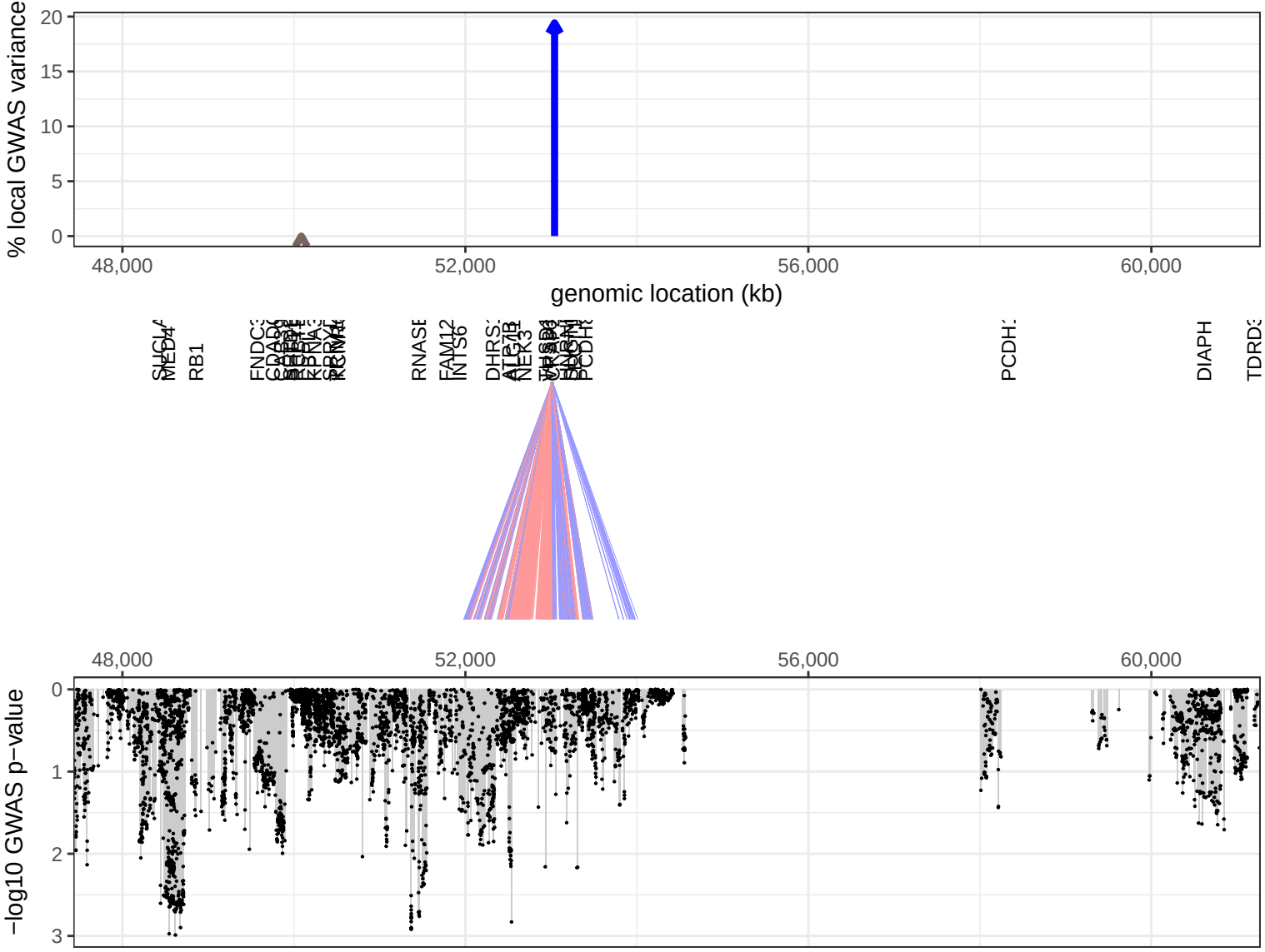
% local GWAS variance
20
15
10
5
0
48,000
52,000
56,000
60,000
genomic location (kb)
MED4
RB1
RNASE
FAM12
INTS6
PCDH
DIAPH
TDRD
48,000
52,000
56,000
60,000
−log10 GWAS p−value
0
1
2
3