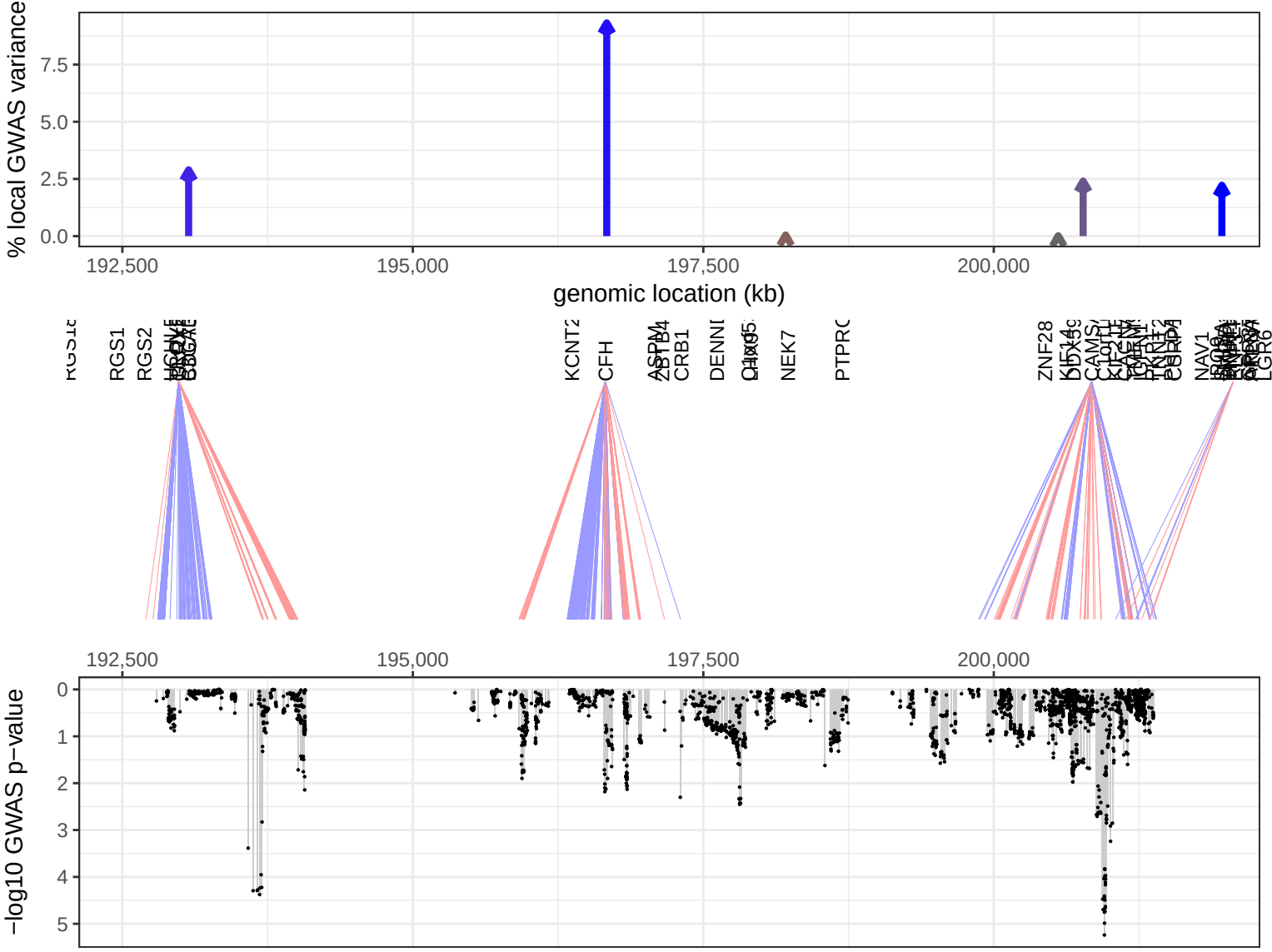
% local GWAS variance
7.5
5.0
2.5
0.0
192,500
195,000
197,500
200,000
genomic location (kb)
RGS1
RGS2
KCNT2
CFH
CRB1
DENNI
NEK7
PTPRC
ZNF28
NAV1
192,500
195,000
197,500
200,000
−log10 GWAS p−value
0
1
2
3
4
5