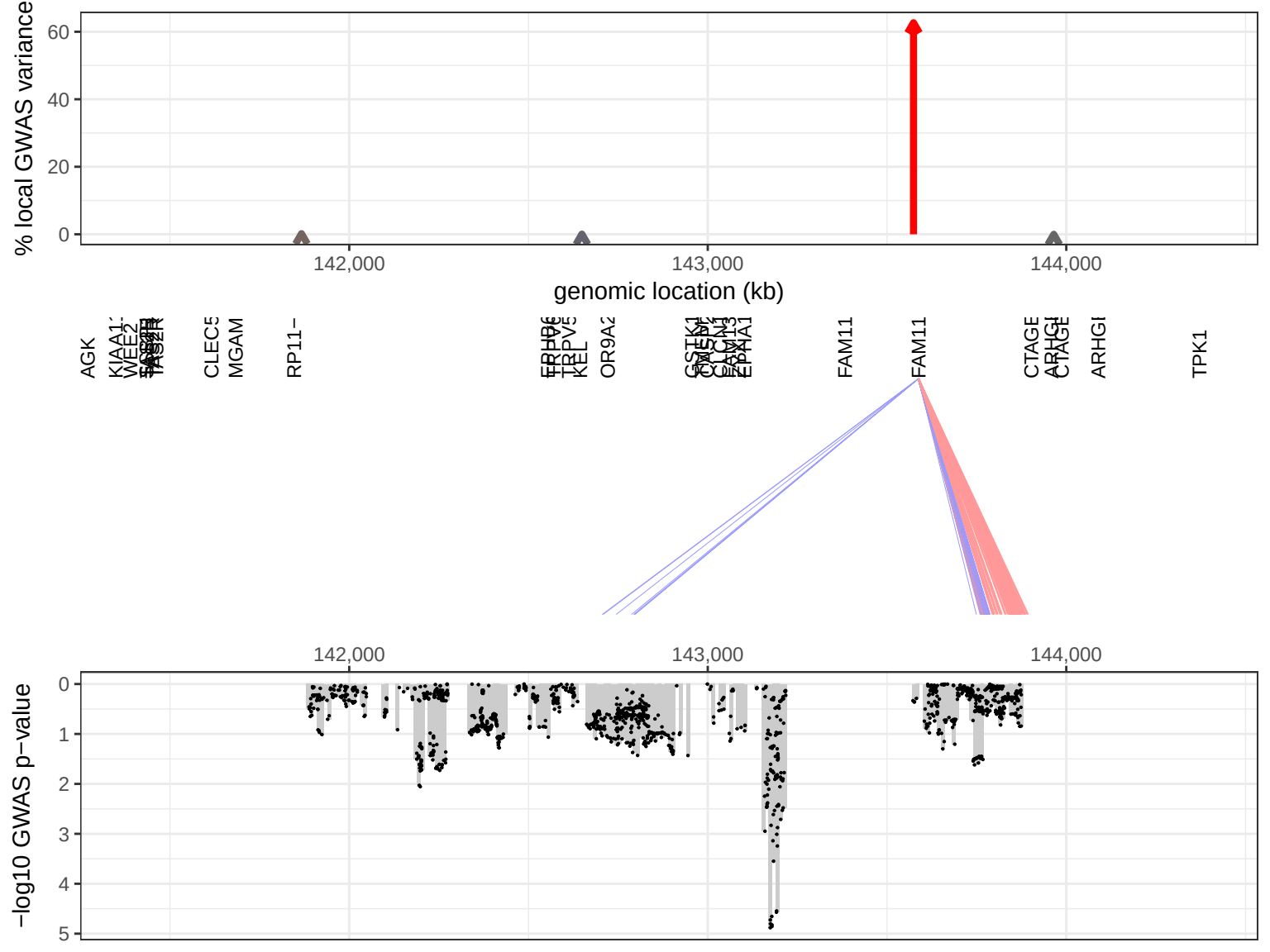
% local GWAS variance
60
40
20
0
142,000
143,000
144,000
genomic location (kb)
AGK
CLEC5
MGAM
RP11−
KEL
OR9A2
FAM11
FAM11
CTAGE
ARHGI
TPK1
142,000
143,000
144,000
−log10 GWAS p−value
0
1
2
3
4
5