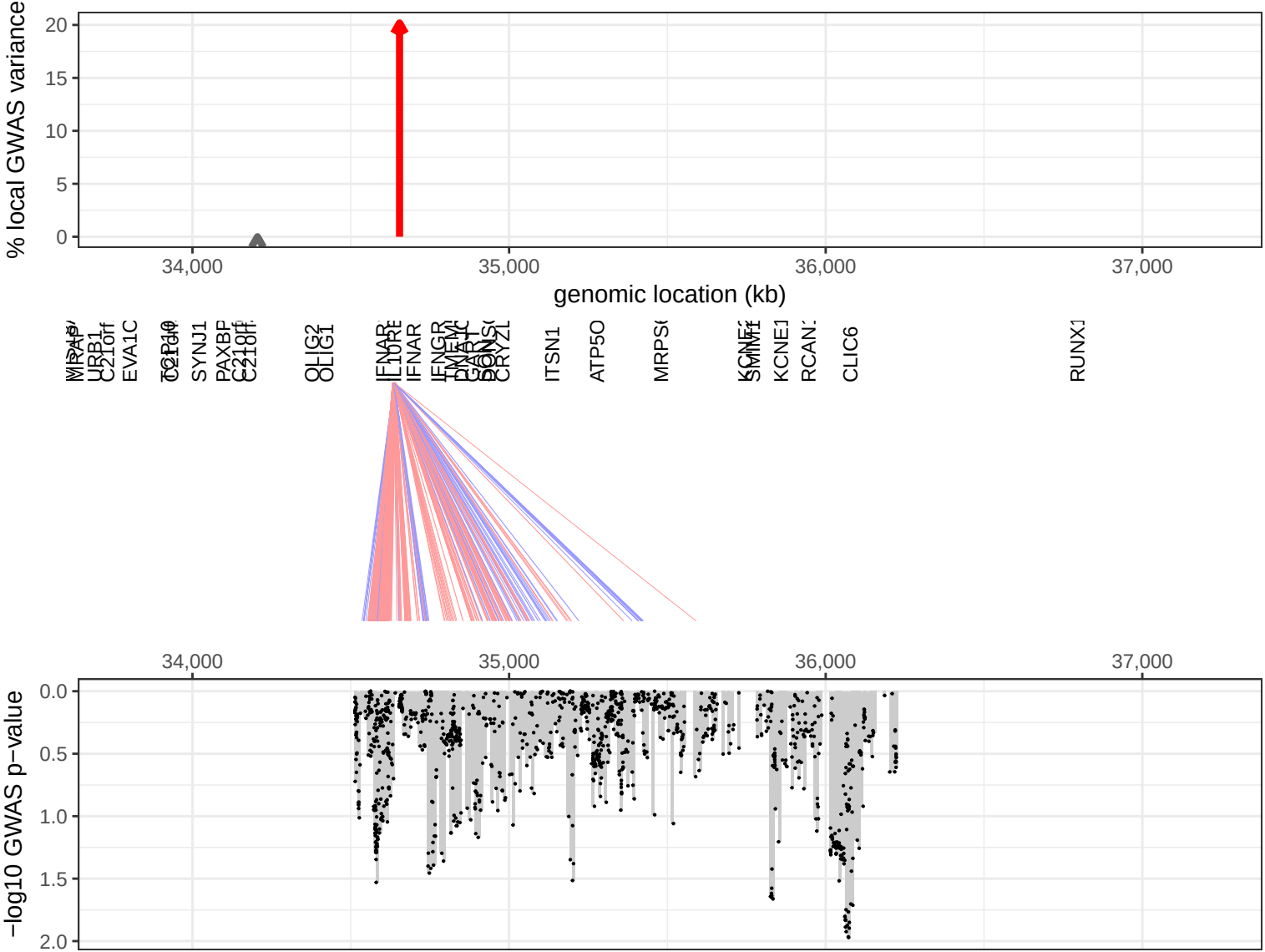
% local GWAS variance
20
15
10
5
0
34,000
35,000
36,000
37,000
genomic location (kb)
EVA1C
SYNJ1
PAXBP
OLIG2
OLIG1
IFNAR
IFNAR
ITSN1
ATP5O
MRPS
KCNE1
RCAN
CLIC6
RUNX1
34,000
35,000
36,000
37,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0