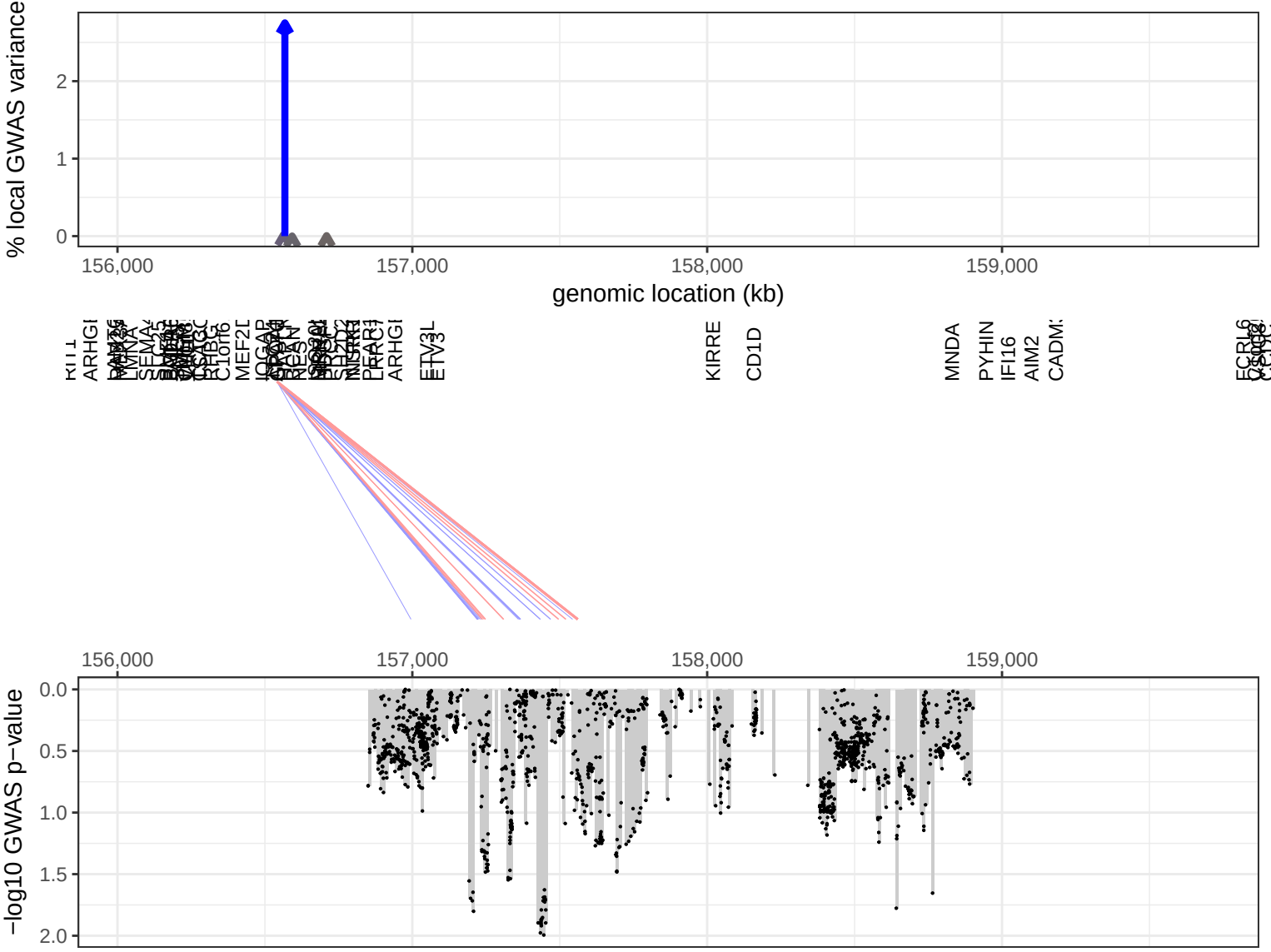
% local GWAS variance
2
1
0
156,000
157,000
158,000
159,000
genomic location (kb)
ARHGI
ETV3L
ETV3
KIRRE
CD1D
MNDA
PYHIN
IFI16
AIM2
CADM
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0