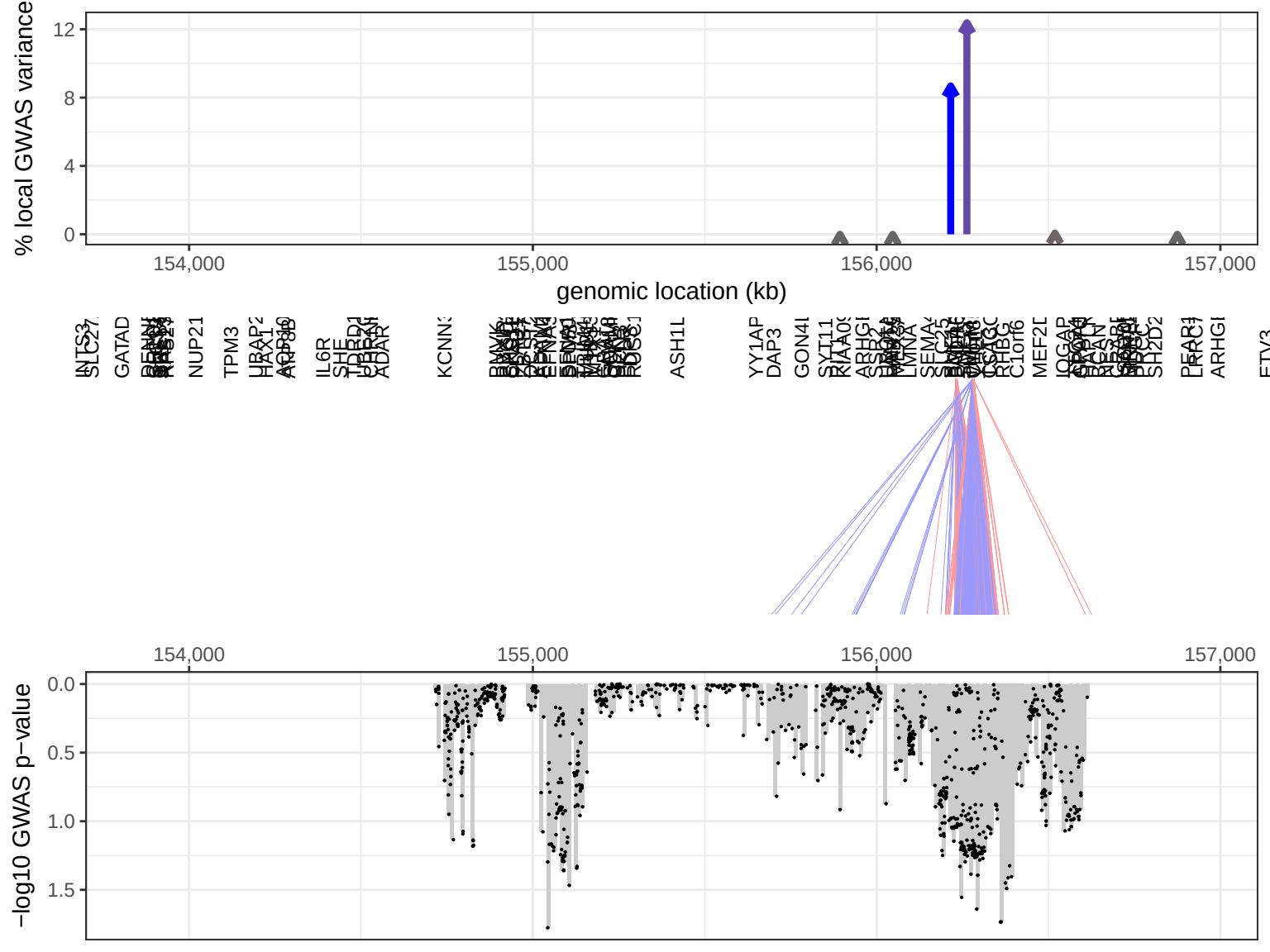
% local GWAS variance
12
8
4
0
154,000
155,000
156,000
157,000
genomic location (kb)
GATAD
NUP21
TPM3
IL6R
KCNN3
ASH1L
YY1AP
DAP3
GON4L
MEF2D
ARHGI
ETV3
154,000
155,000
156,000
157,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5