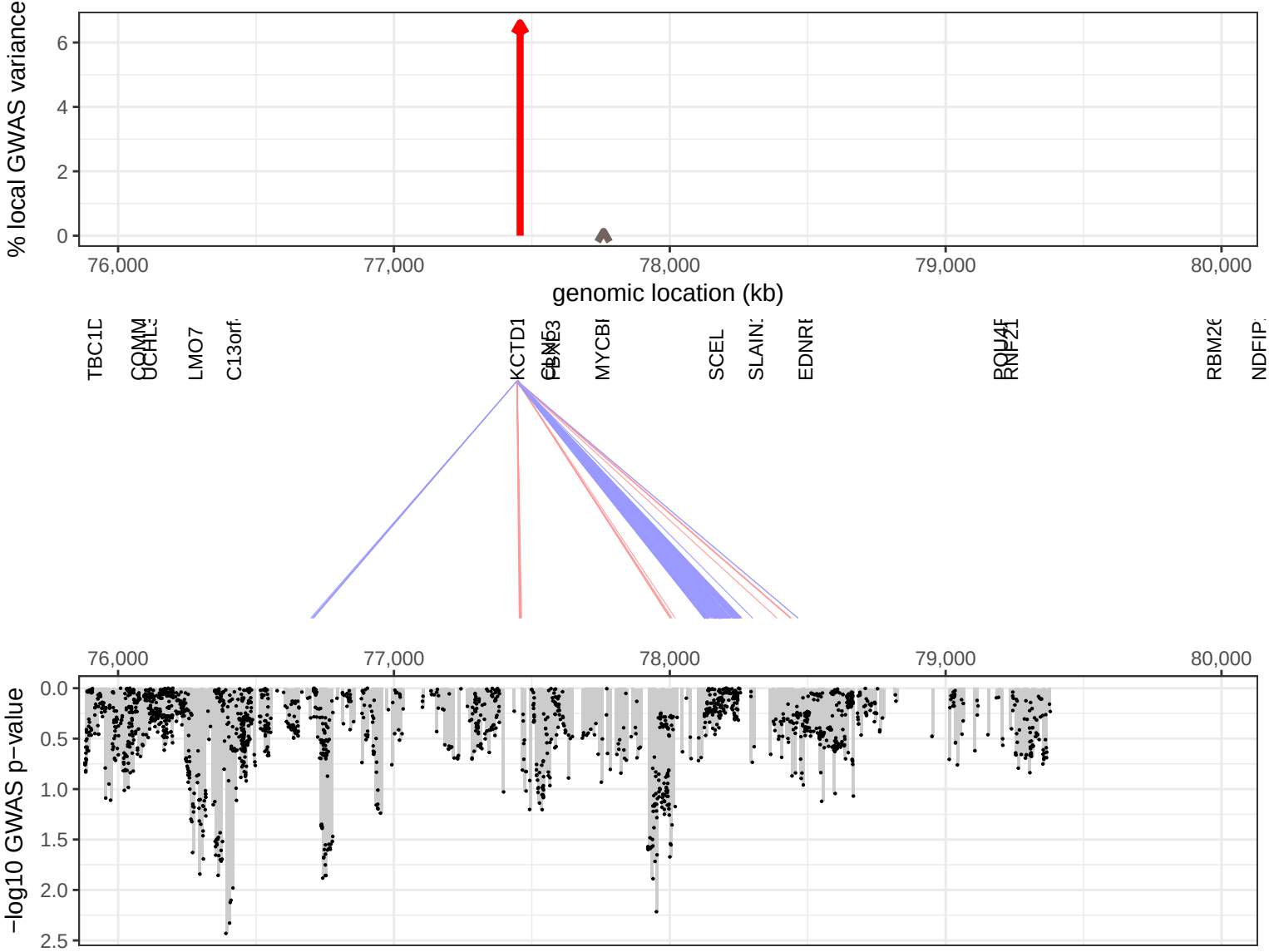
% local GWAS variance
6
4
2
0
76,000
77,000
78,000
79,000
80,000
genomic location (kb)
LMO7
C13orf
KCTD1
MYCBI
SCEL
SLAIN
EDNRI
RBM26
NDFIP
76,000
77,000
78,000
79,000
80,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5