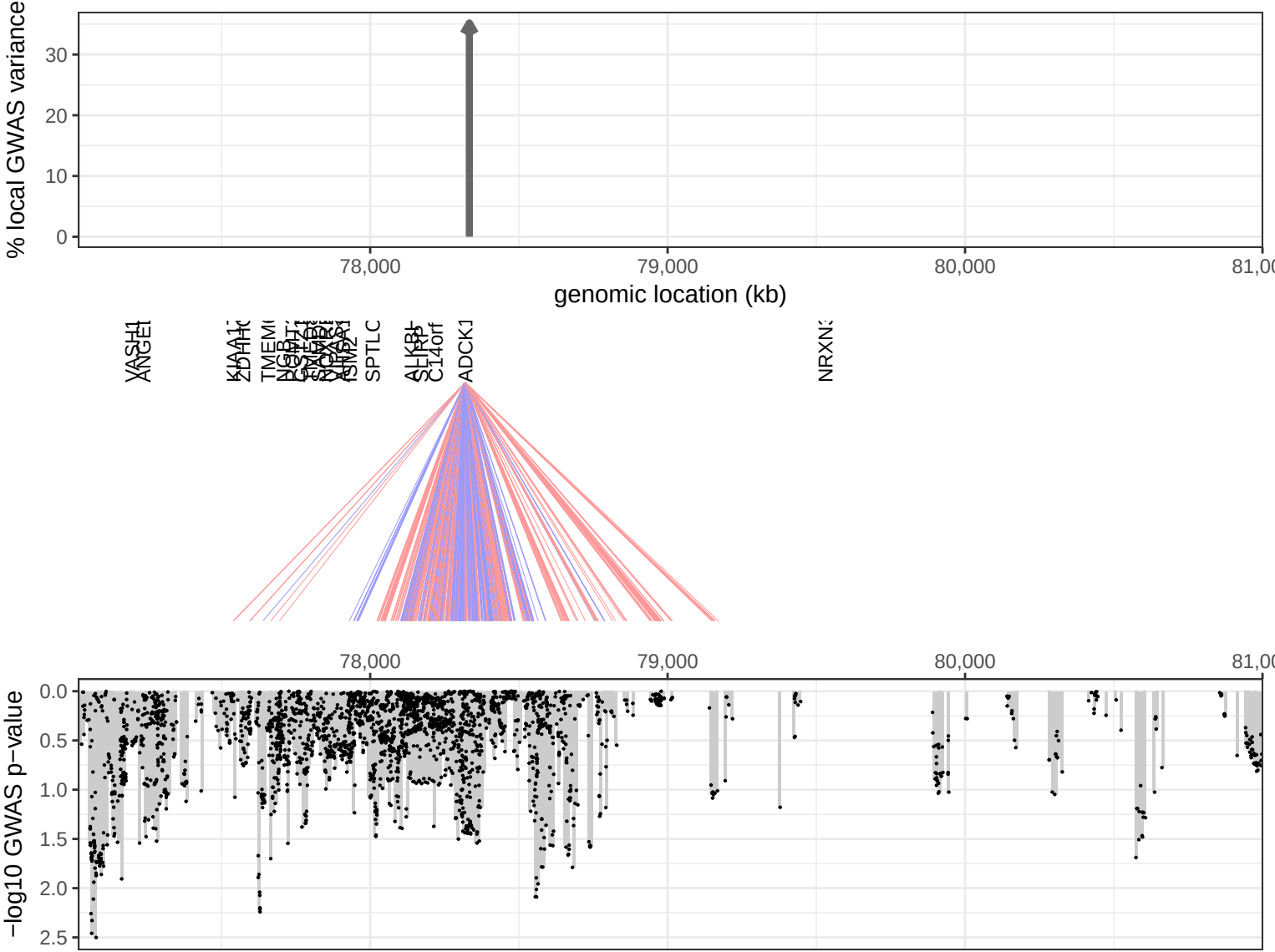
% local GWAS variance
30
20
10
0
78,000
79,000
80,000
genomic location (kb)
SPTLC
ADCK1
NRXN
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5