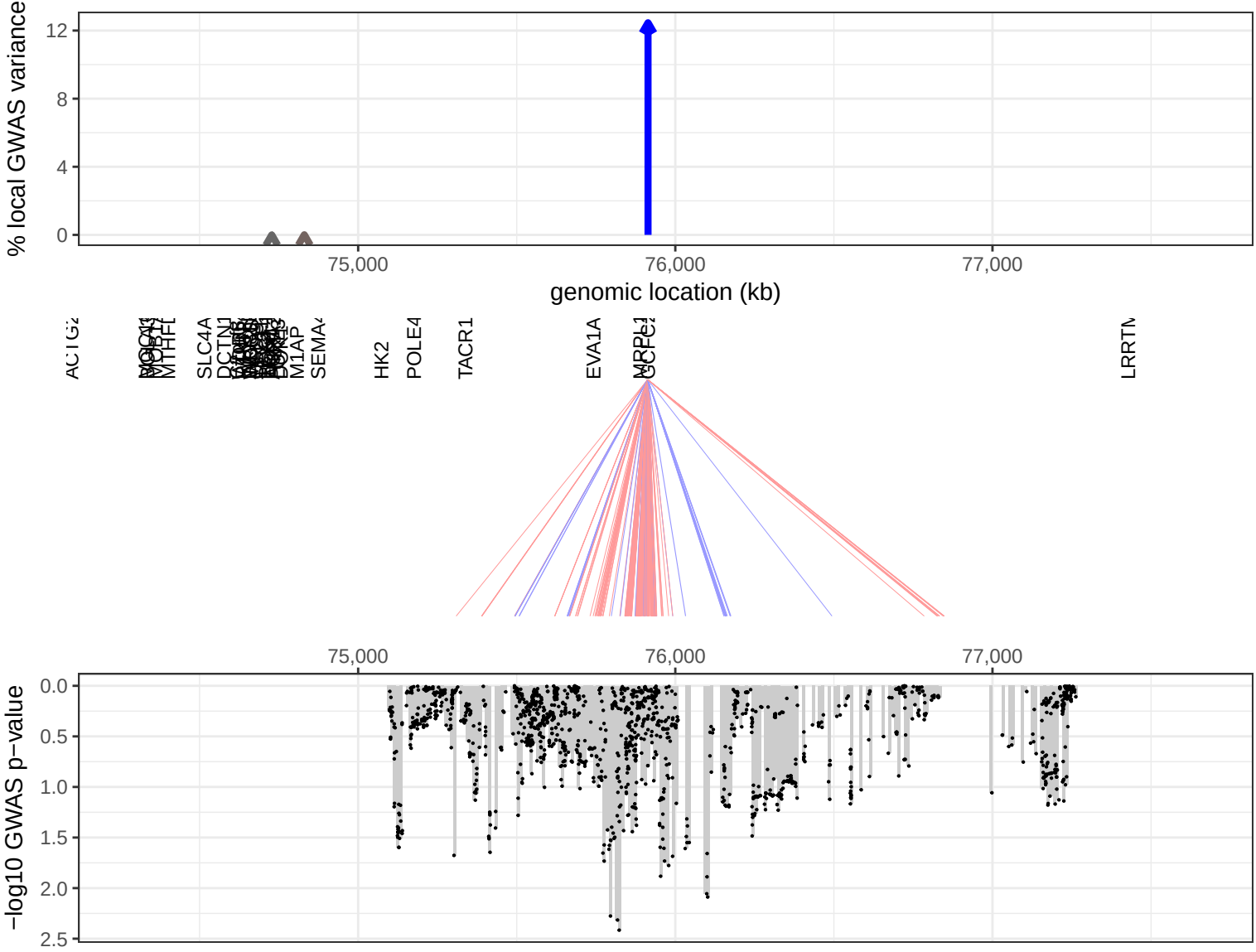
% local GWAS variance
12
8
4
0
75,000
76,000
77,000
genomic location (kb)
SLC4A
HK2
POLE4
TACR1
EVA1A
M1AP
SEMA
LRRTM
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5