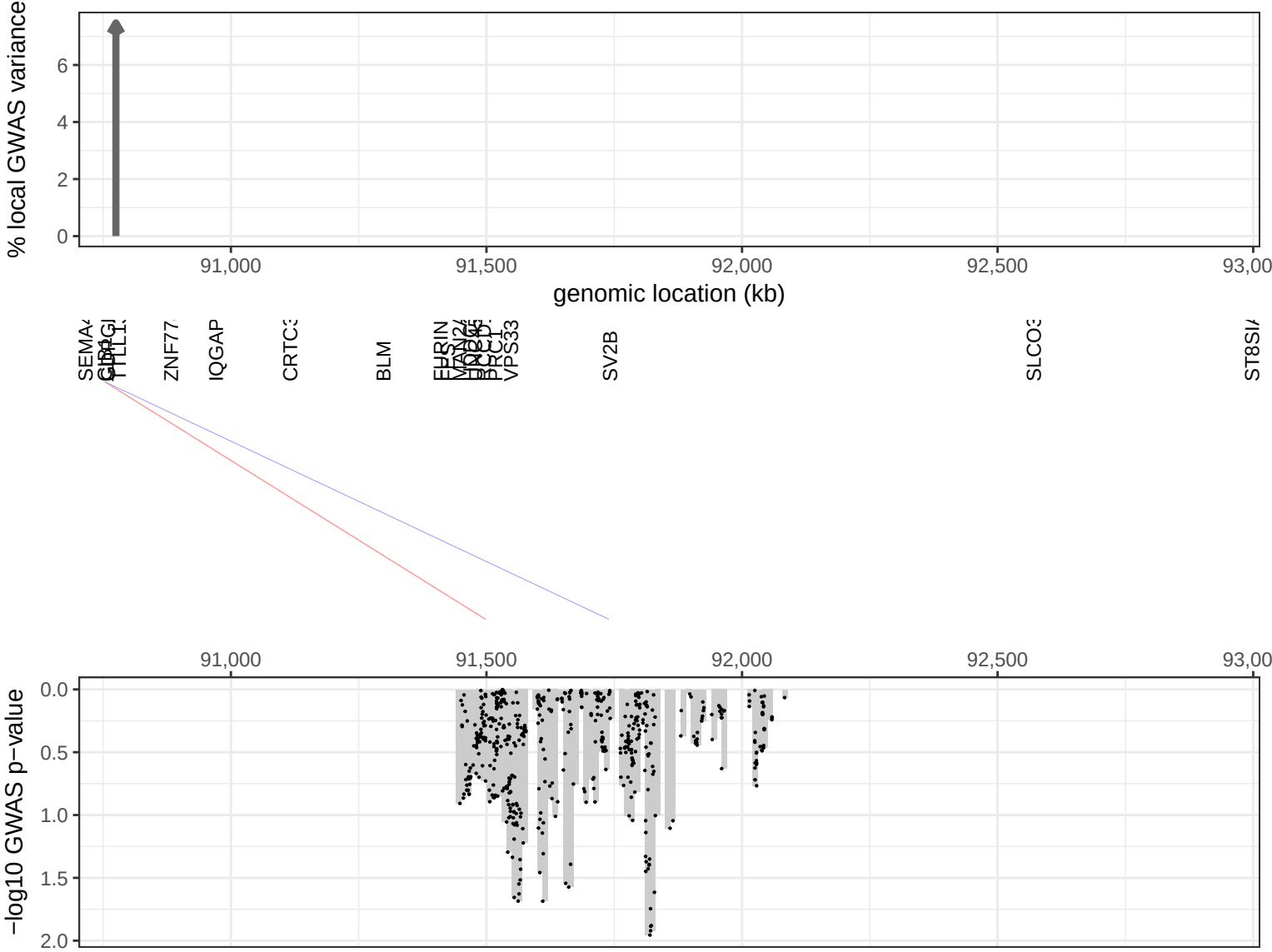
% local GWAS variance
0
2
4
6
91,000
91,500
92,000
92,500
93,00
genomic location (kb)
ZNF77
IQGAP
CRTC
BLM
VPS33
SV2B
SLCO
ST8SI
91,000
91,500
92,000
92,500
93,00
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0