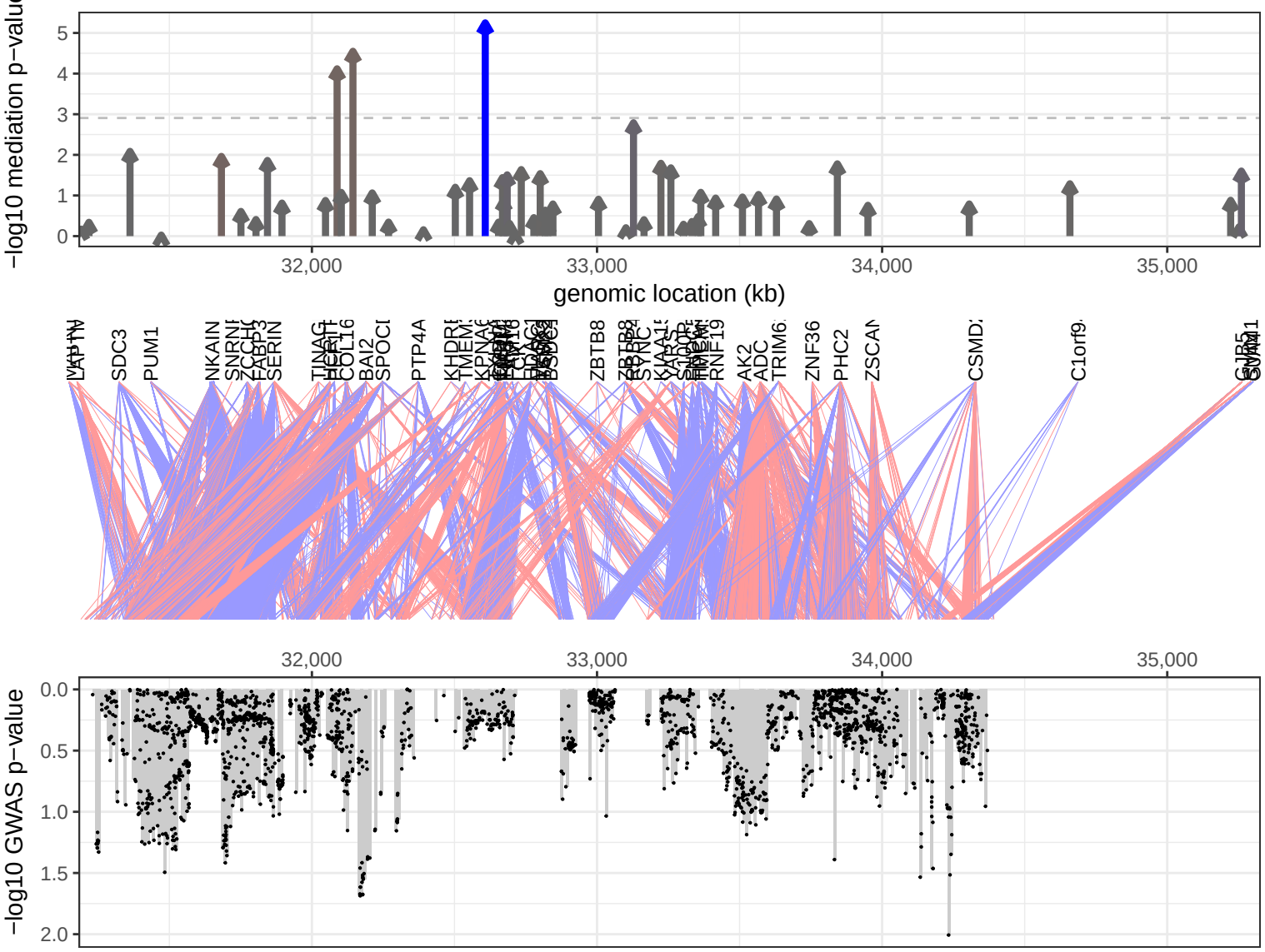
−log10 mediation p−value
5
4
3
2
1
0
32,000
33,000
34,000
35,000
genomic location (kb)
SDC3
PUM1
NKAIN
SERIN
BAI2
PTP4A
ZBTB8
AK2
ADC
ZNF36
PHC2
CSMD
C1orf9
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0