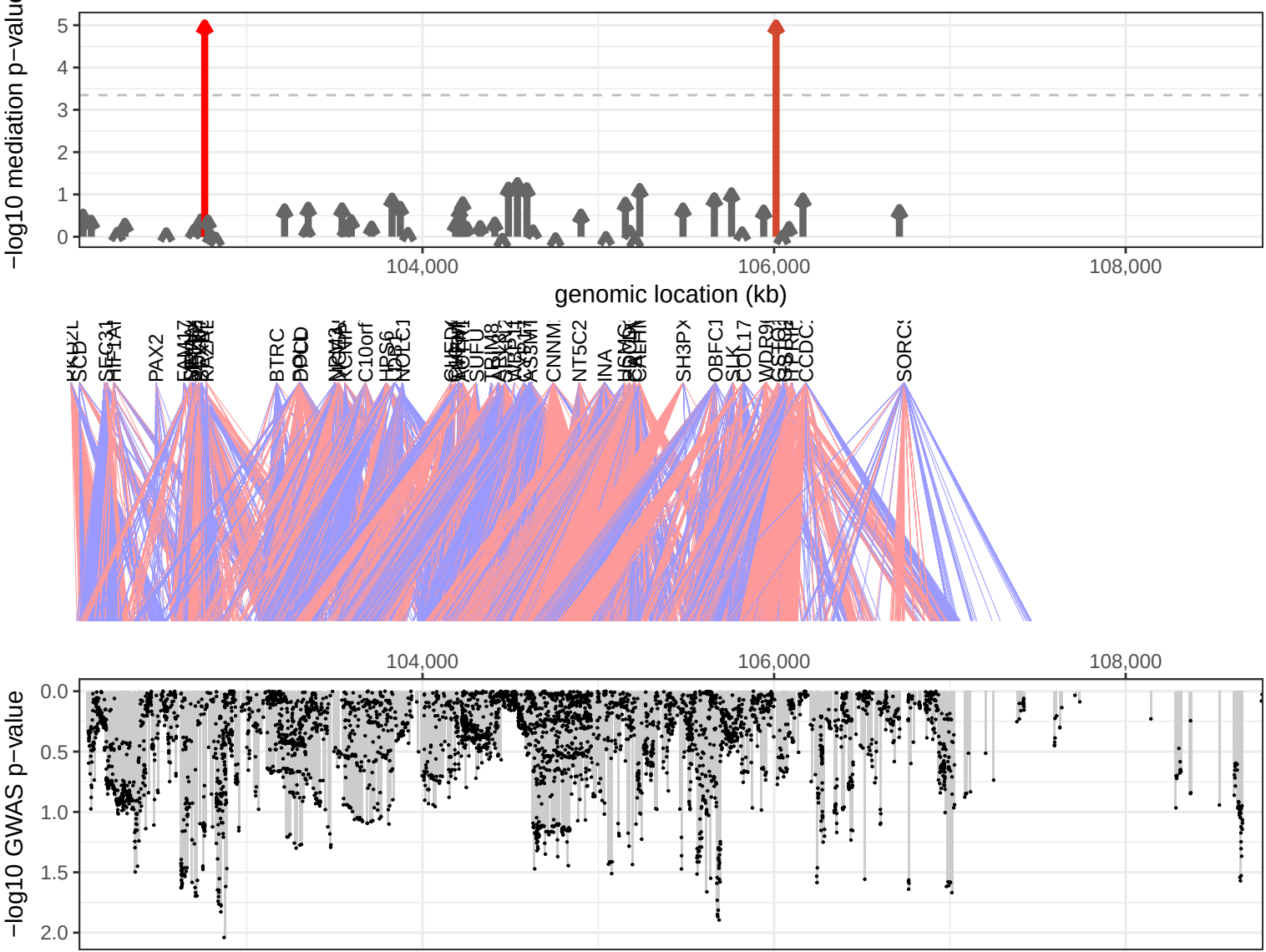
−log10 mediation p−value
5
4
3
2
1
0
104,000
106,000
108,000
genomic location (kb)
PAX2
BTRC
C10orf
CNNM
NT5C2
INA
SH3PX
OBFC1
SORC
104,000
106,000
108,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0