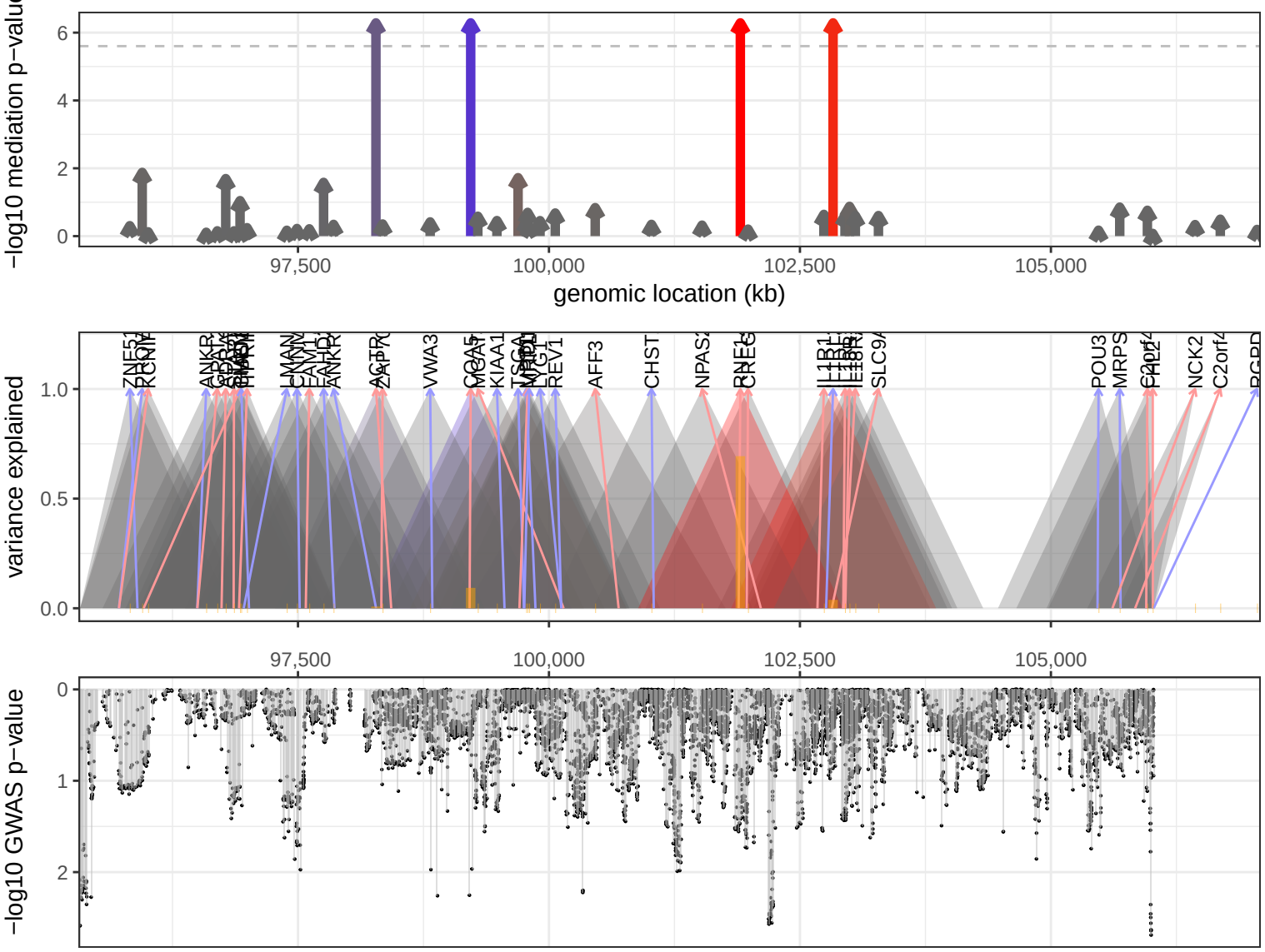
−log10 mediation p−value
6
4
2
0
97,500
100,000
102,500
105,000
genomic location (kb)
variance explained
1.0
0.5
0.0
VWA3
KIAA1
REV1
AFF3
CHST
POU3
MRPS
NCK2
C2orf4
97,500
100,000
102,500
105,000
−log10 GWAS p−value
0
1
2