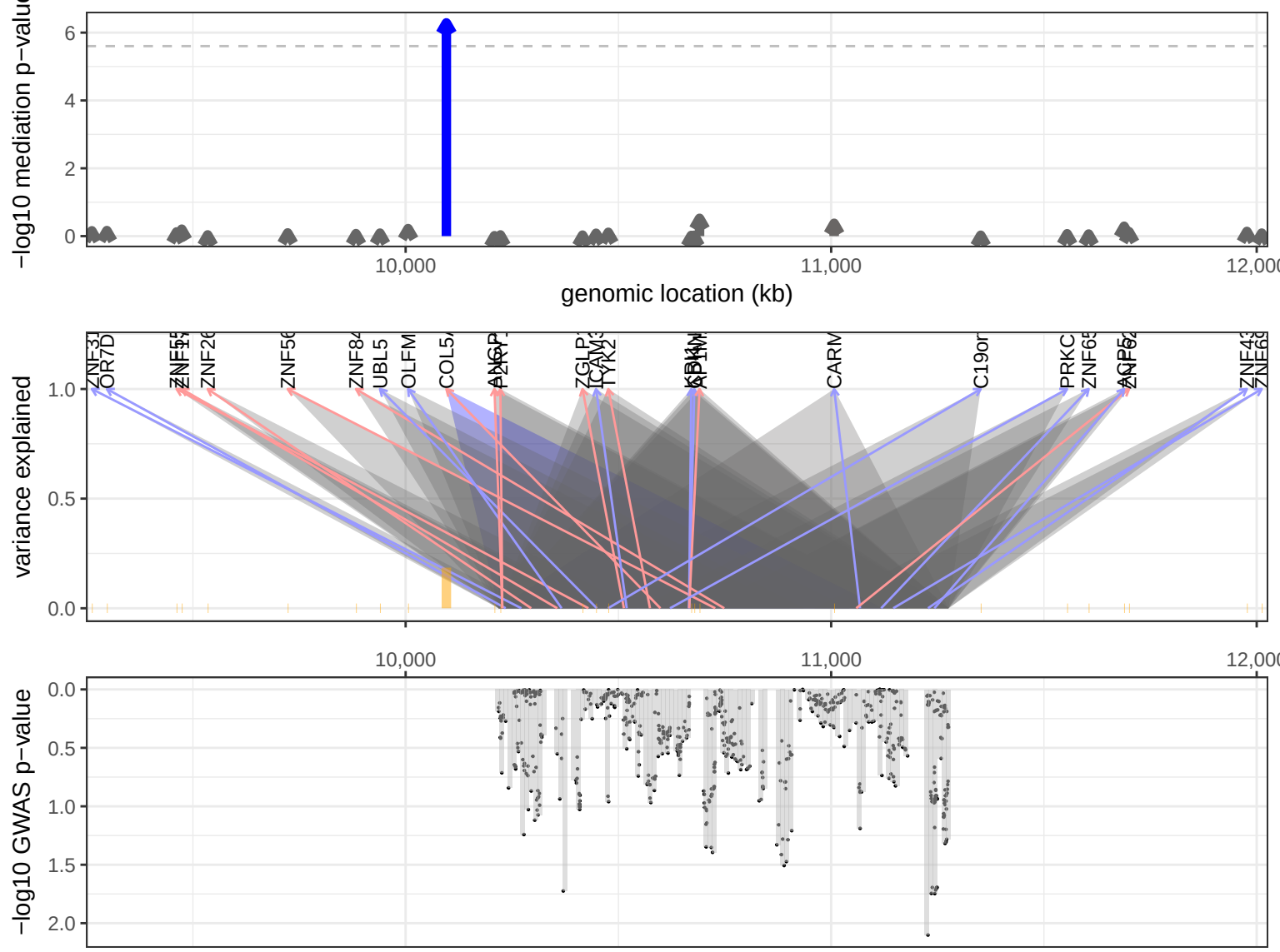
−log10 mediation p−value
6
4
2
0
10,000
11,000
12,000
genomic location (kb)
variance explained
1.0
0.5
0.0
OR7D
ZNF26
ZNF56
ZNF84
UBL5
OLFM
CARM
C19or
PRKC
ZNF65
10,000
11,000
12,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0