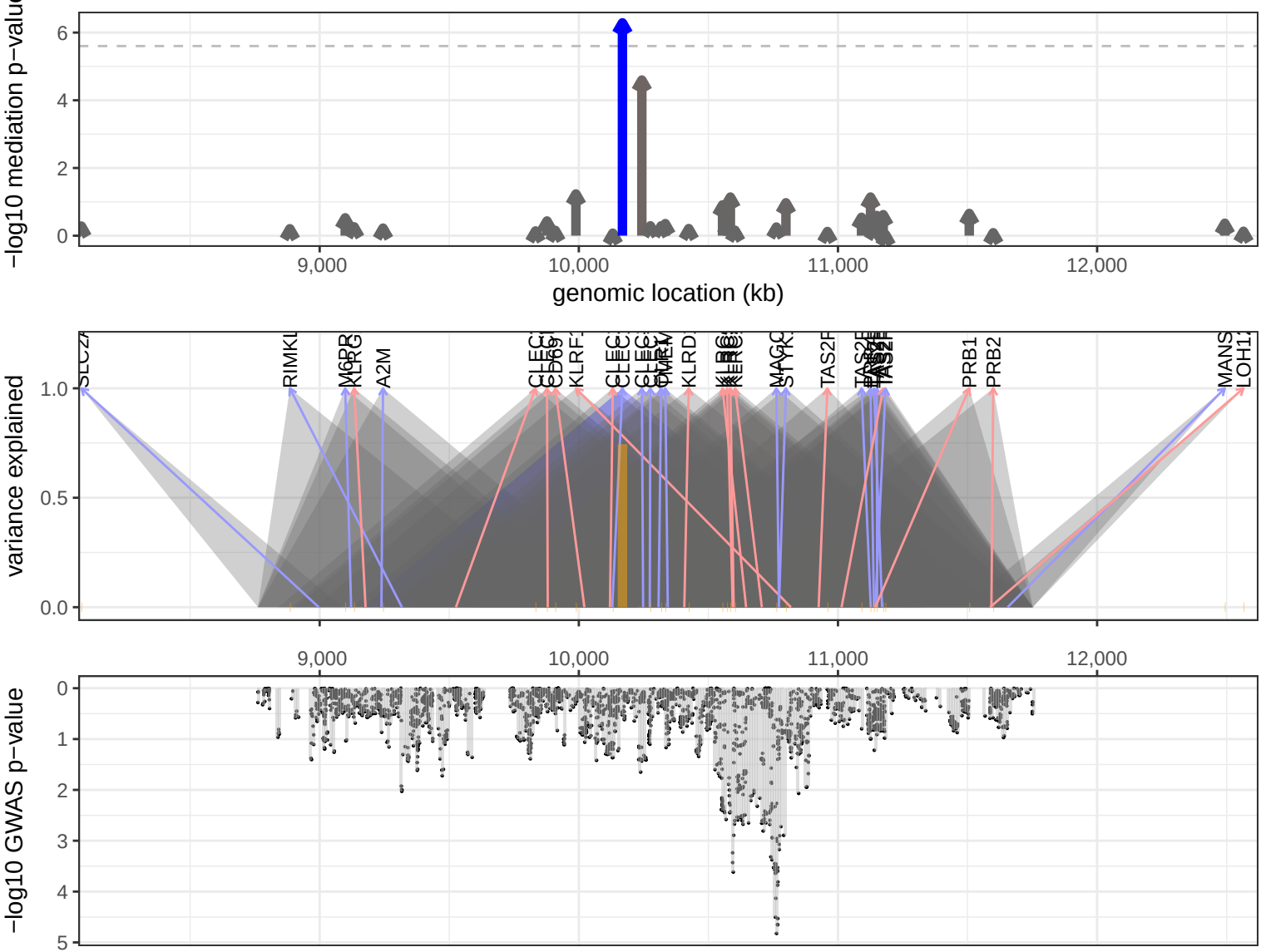
−log10 mediation p−value
6
4
2
0
9,000
10,000
11,000
12,000
genomic location (kb)
variance explained
1.0
0.5
0.0
RIMKL
A2M
KLRD
TAS2F
PRB1
PRB2
MANS
9,000
10,000
11,000
12,000
−log10 GWAS p−value
0
1
2
3
4
5