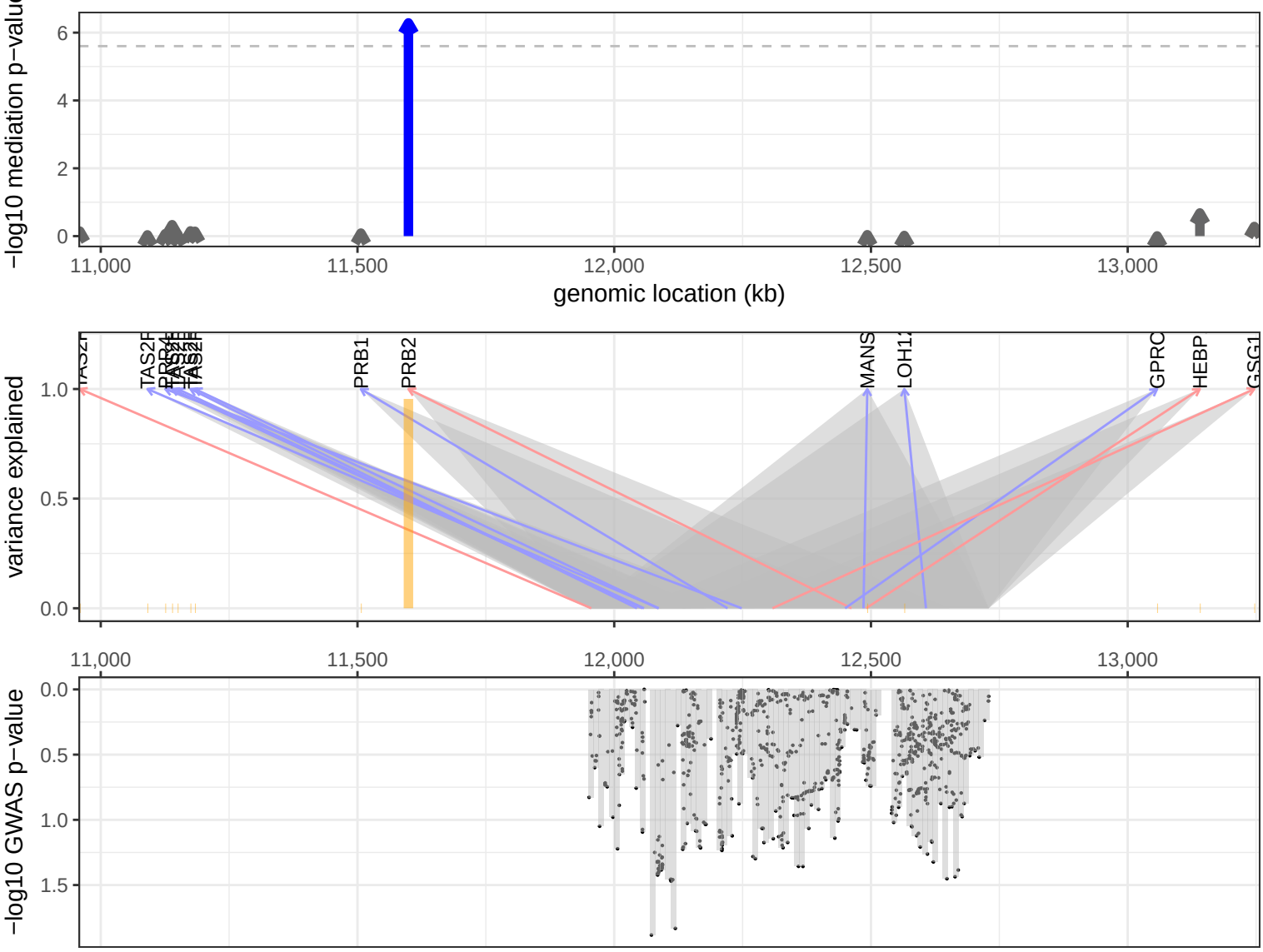
−log10 mediation p−value
6
4
2
0
11,000
11,500
12,000
12,500
13,000
genomic location (kb)
variance explained
1.0
0.5
0.0
TAS2F
PRB1
PRB2
MANS
LOH1
GPRC
HEBP
GSG1
−log10 GWAS p−value
0.0
0.5
1.0
1.5