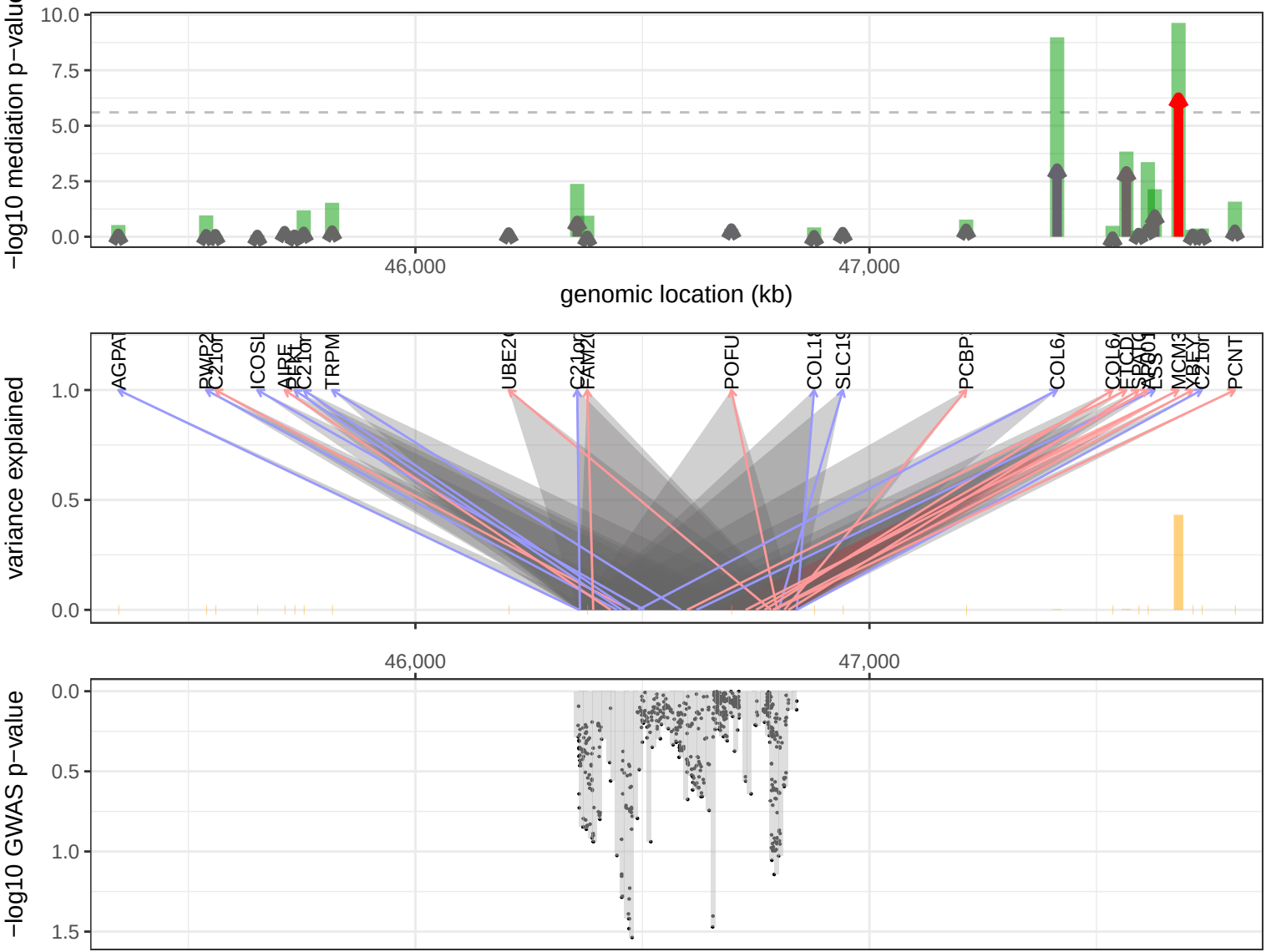
−log10 mediation p−value
10.0
7.5
5.0
2.5
0.0
46,000
47,000
genomic location (kb)
variance explained
1.0
0.5
0.0
AGPAT
TRPM
UBE2
POFU
COL19
SLC19
PCBP
COL6
PCNT
46,000
47,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5