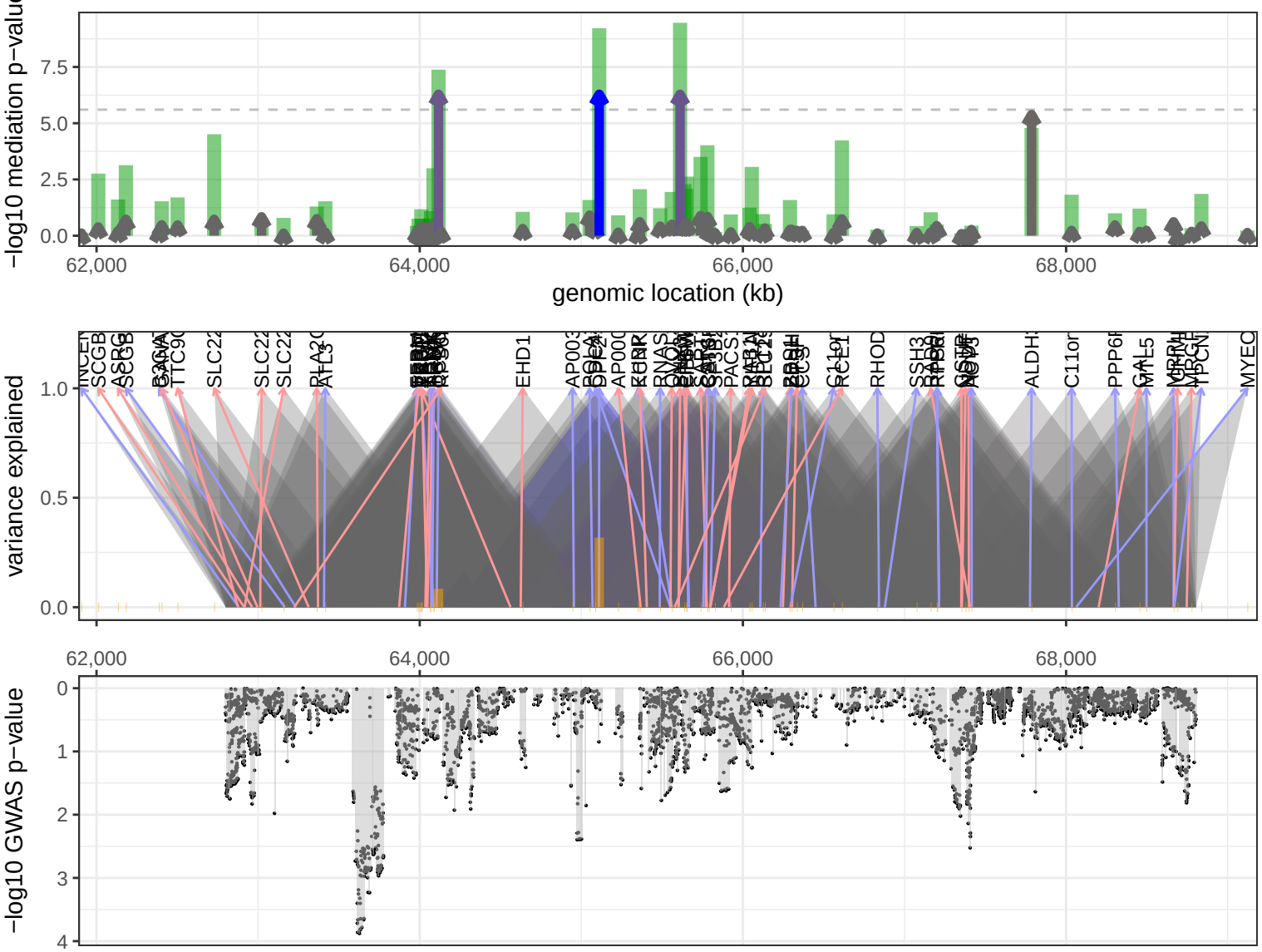
−log10 mediation p−value
7.5
5.0
2.5
0.0
62,000
64,000
66,000
68,000
genomic location (kb)
variance explained
1.0
0.5
0.0
EHD1
RHOD
ALDH
MYEO
−log10 GWAS p−value
0
1
2
3
4