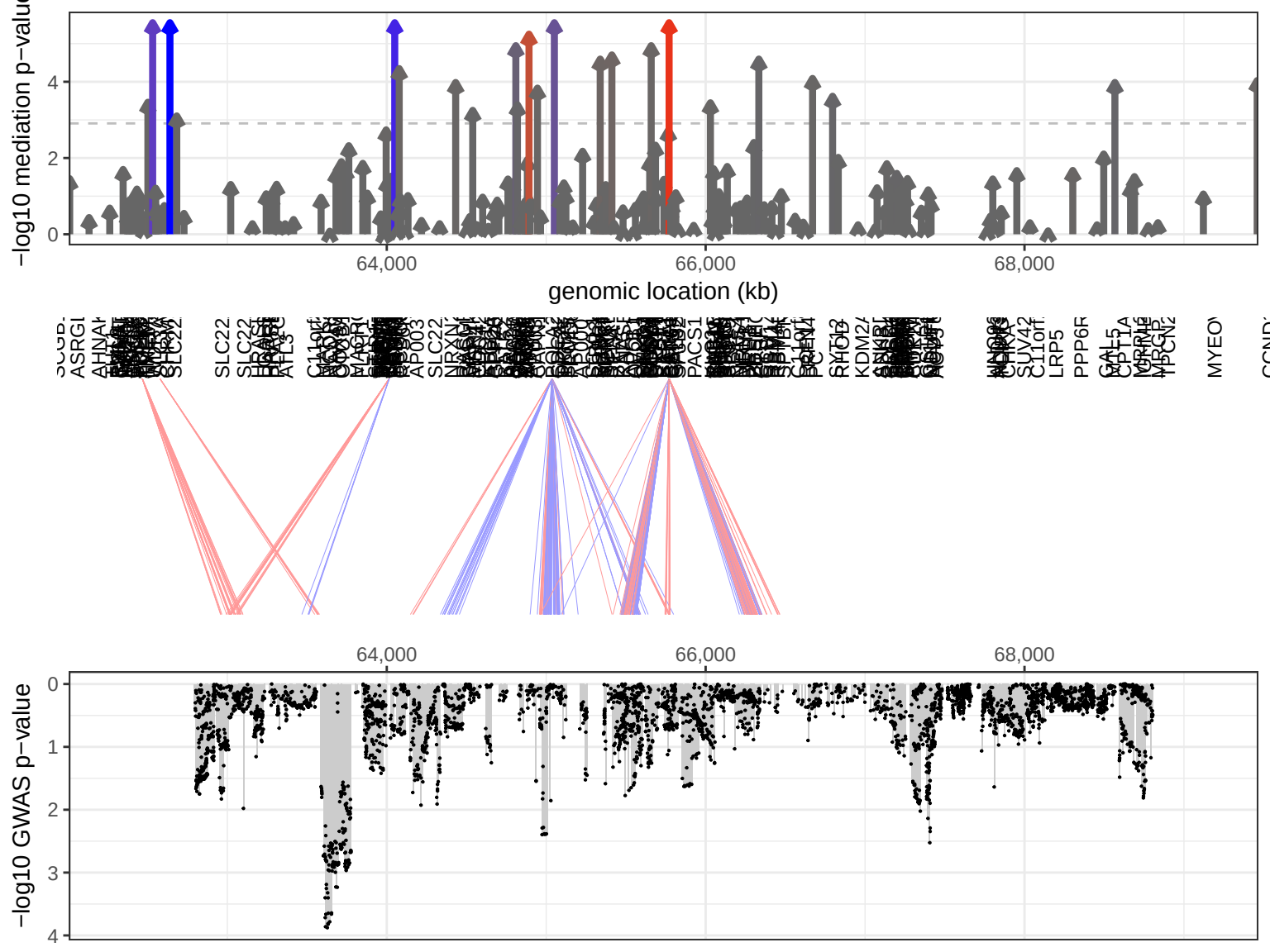
−log10 mediation p−value
0
2
4
64,000
66,000
68,000
genomic location (kb)
SLC22
AP003
PACS1
RYR3
KDM2A
LRP5
PPP6R
MYEOV
−log10 GWAS p−value
0
1
2
3
4
64,000
66,000
68,000