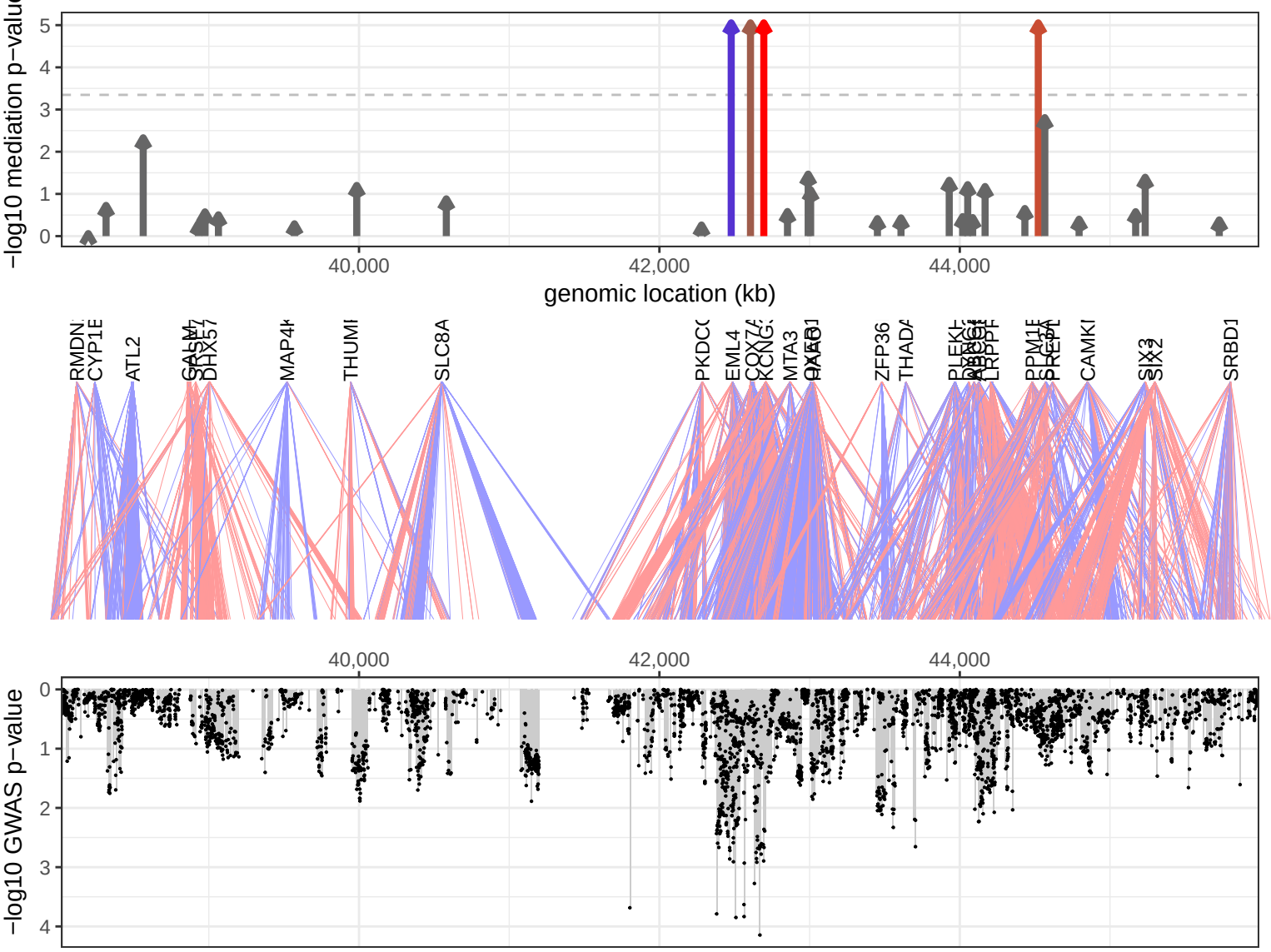
−log10 mediation p−value
5
4
3
2
1
0
40,000
42,000
44,000
genomic location (kb)
RMDN
CYP1E
ATL2
MAP4
THUMI
SLC8A
PKDC
EML4
MTA3
ZFP36
THAD
CAMKI
SIX3
SIX2
SRBD1
40,000
42,000
44,000
−log10 GWAS p−value
0
1
2
3
4