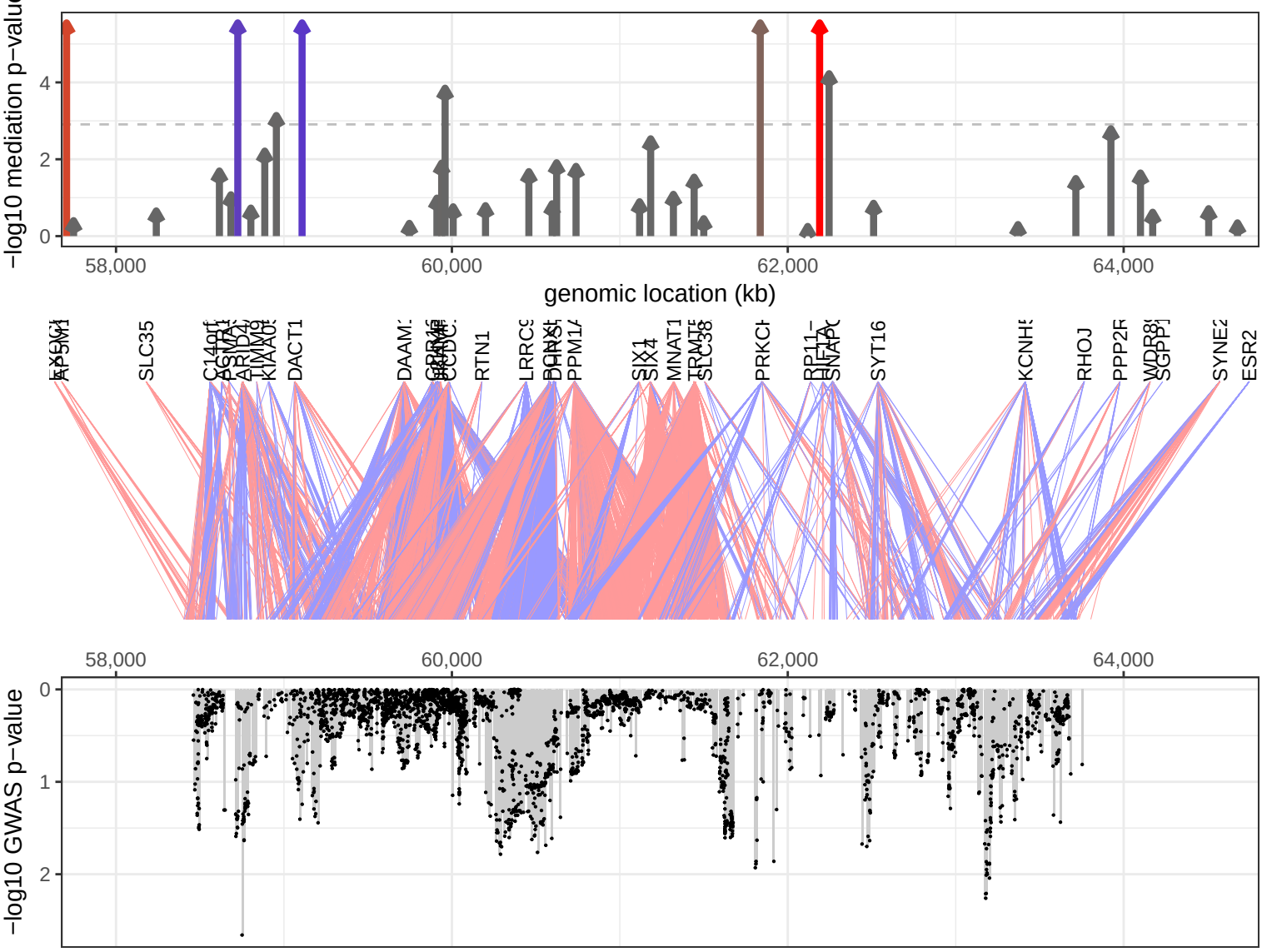
−log10 mediation p−value
0
2
4
58,000
60,000
62,000
64,000
genomic location (kb)
SLC35
DACT1
RTN1
MNAT1
PRKCH
SYT16
KCNH
RHOJ
PPP2R
SYNE2
ESR2
−log10 GWAS p−value
0
1
2