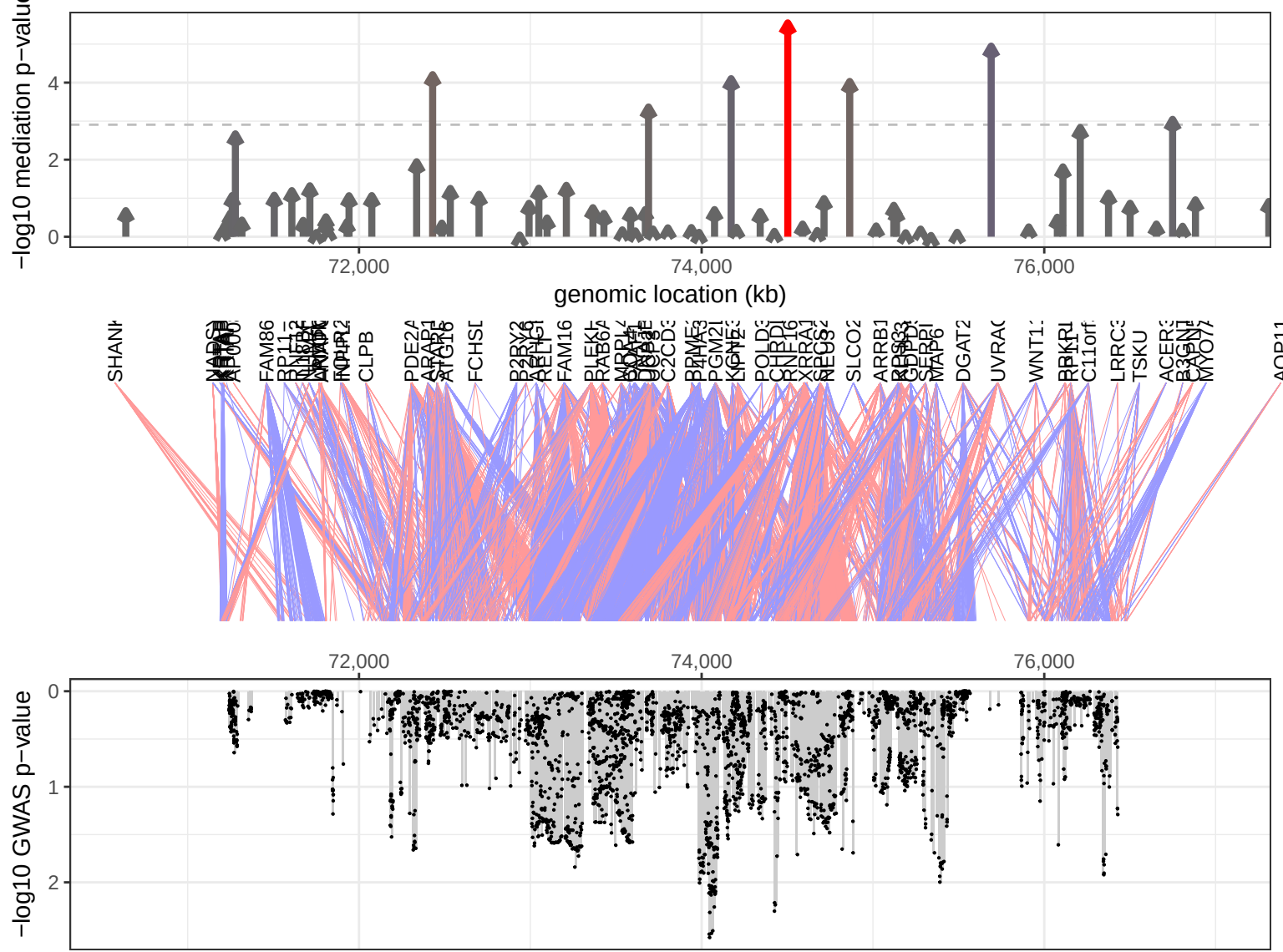
−log10 mediation p−value
0
2
4
72,000
74,000
76,000
genomic location (kb)
SHANK
FAM86
CLPB
PDE2A
FCHSD
FAM16
SLCO2
ARRB1
DGAT2
UVRAG
WNT11
C11orf
LRRC
TSKU
MYO7A
−log10 GWAS p−value
0
1
2
72,000
74,000
76,000