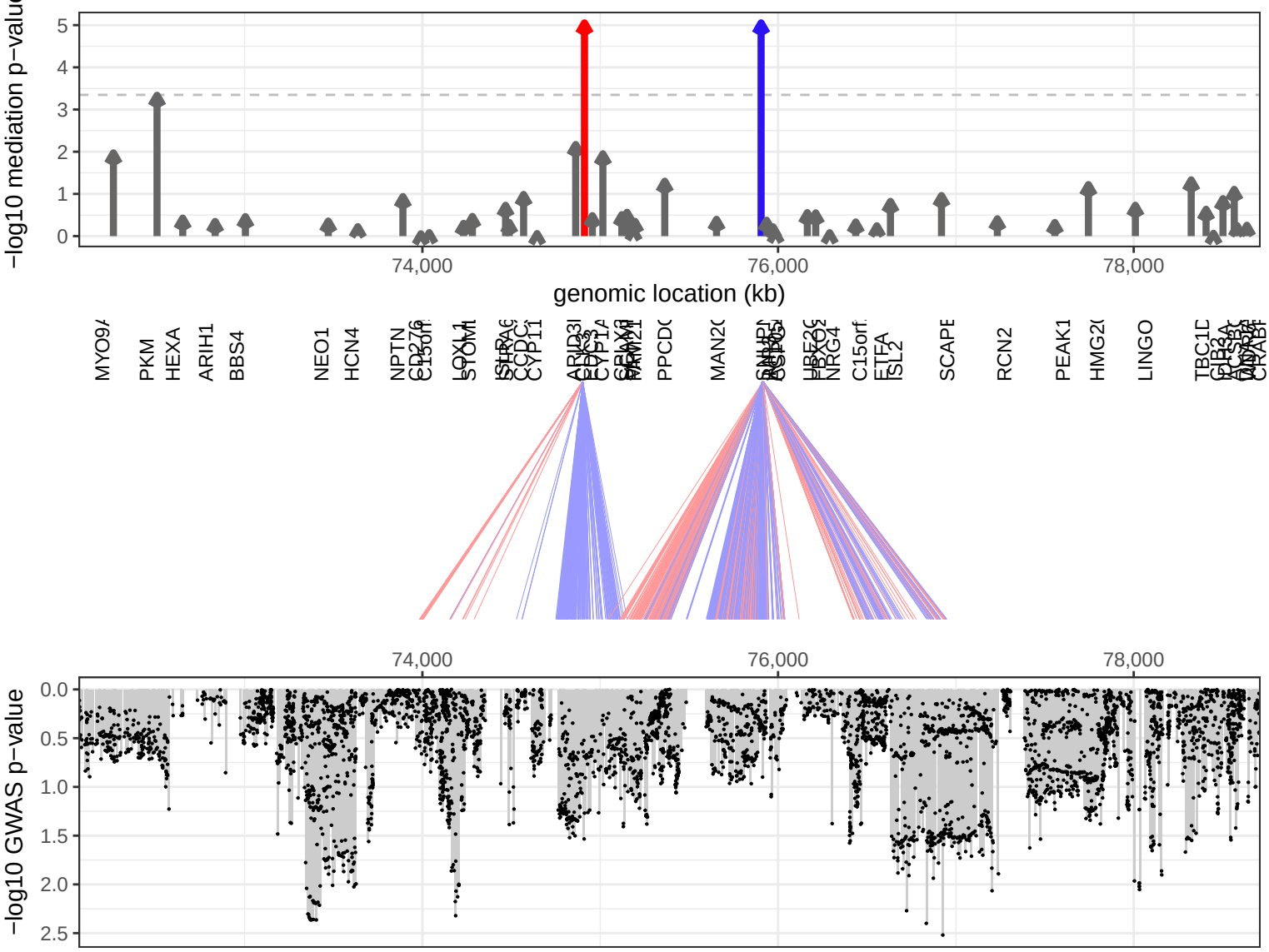
−log10 mediation p−value
5
4
3
2
1
0
74,000
76,000
78,000
genomic location (kb)
MYO9A
PKM
HEXA
ARIH1
BBS4
NEO1
HCN4
NPTN
C15orf
LOXL1
STOM
CYP11
ARID3
CYP1A
PPCD
MAN2C
NRG4
C15orf
ETFA
ISL2
SCAPE
RCN2
PEAK1
HMG2
LINGO
TBC1D
74,000
76,000
78,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5