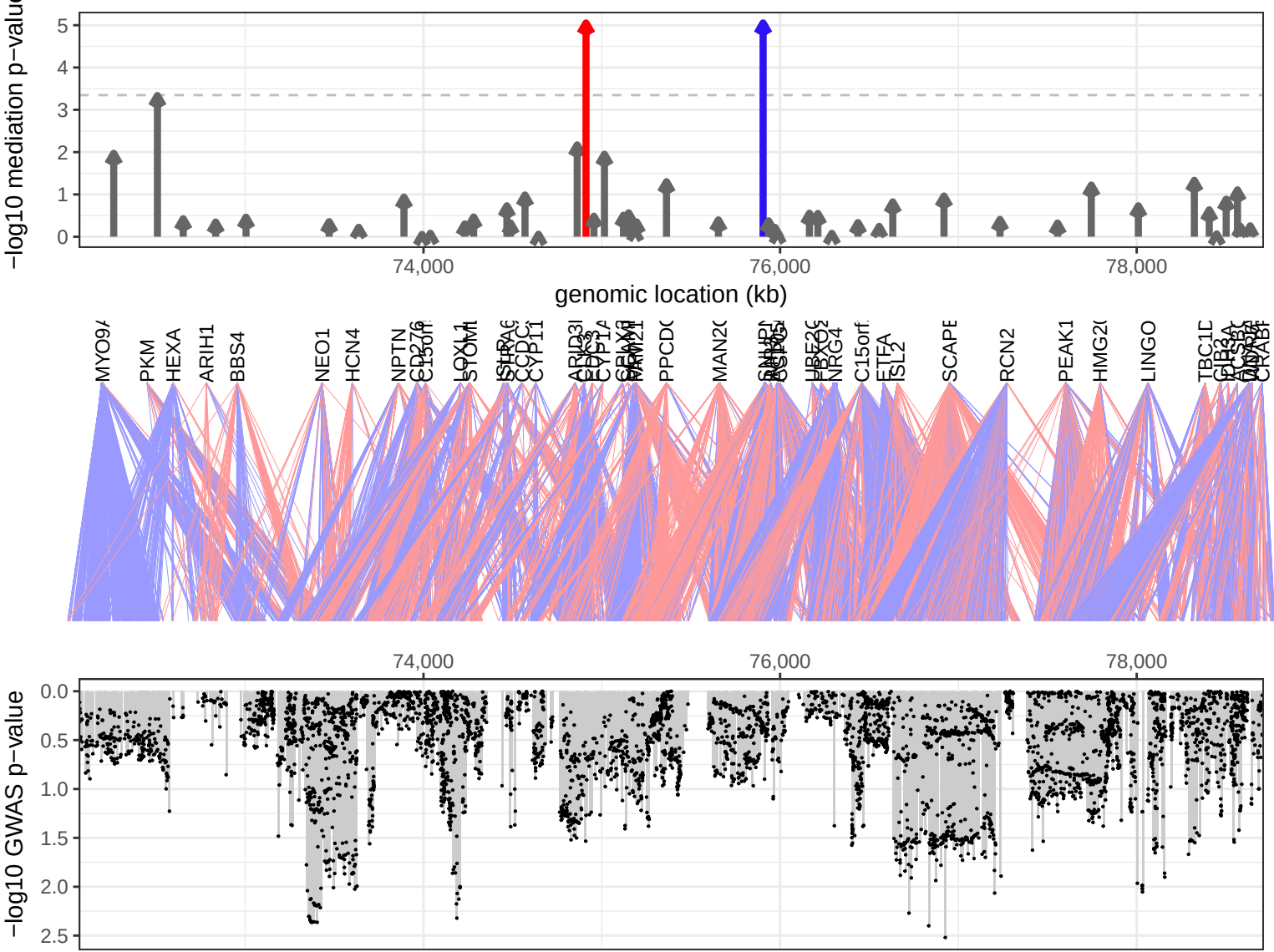
−log10 mediation p−value
5
4
3
2
1
0
74,000
76,000
78,000
genomic location (kb)
PKM
HEXA
ARIH1
BBS4
NEO1
HCN4
NPTN
LOXL1
CYP11
PPCD
MAN2
NRG4
C15orf
ETFA
SCAPE
RCN2
PEAK1
HMG2
LINGO
74,000
76,000
78,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
2.5