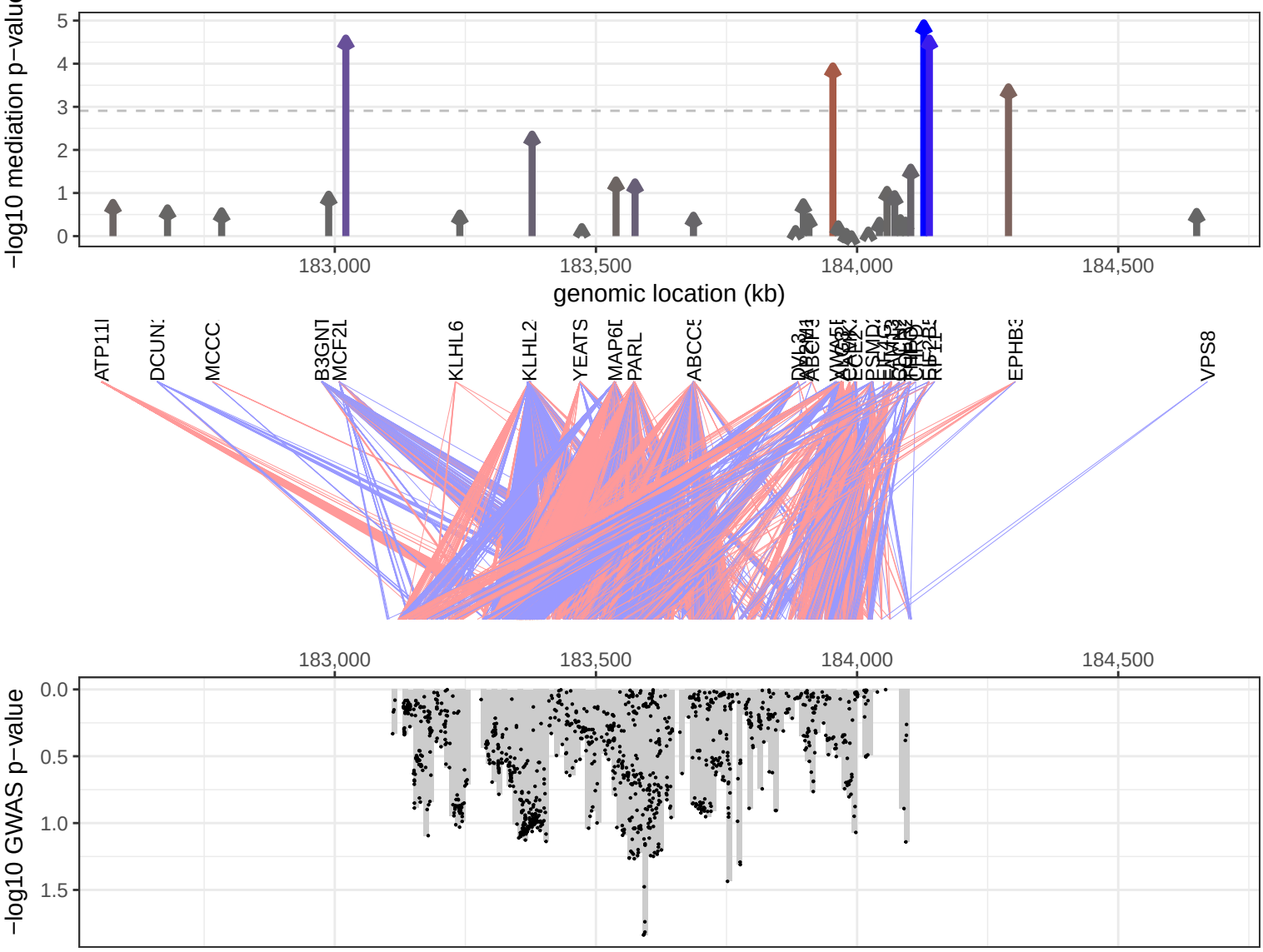
−log10 mediation p−value
5
4
3
2
1
0
183,000
183,500
184,000
184,500
genomic location (kb)
ATP11I
DCUN
MCCC
B3GNT
MCF2L
KLHL6
KLHL2
YEATS
MAP6D
PARL
ABCC5
EPHB3
VPS8
183,000
183,500
184,000
184,500
−log10 GWAS p−value
0.0
0.5
1.0
1.5