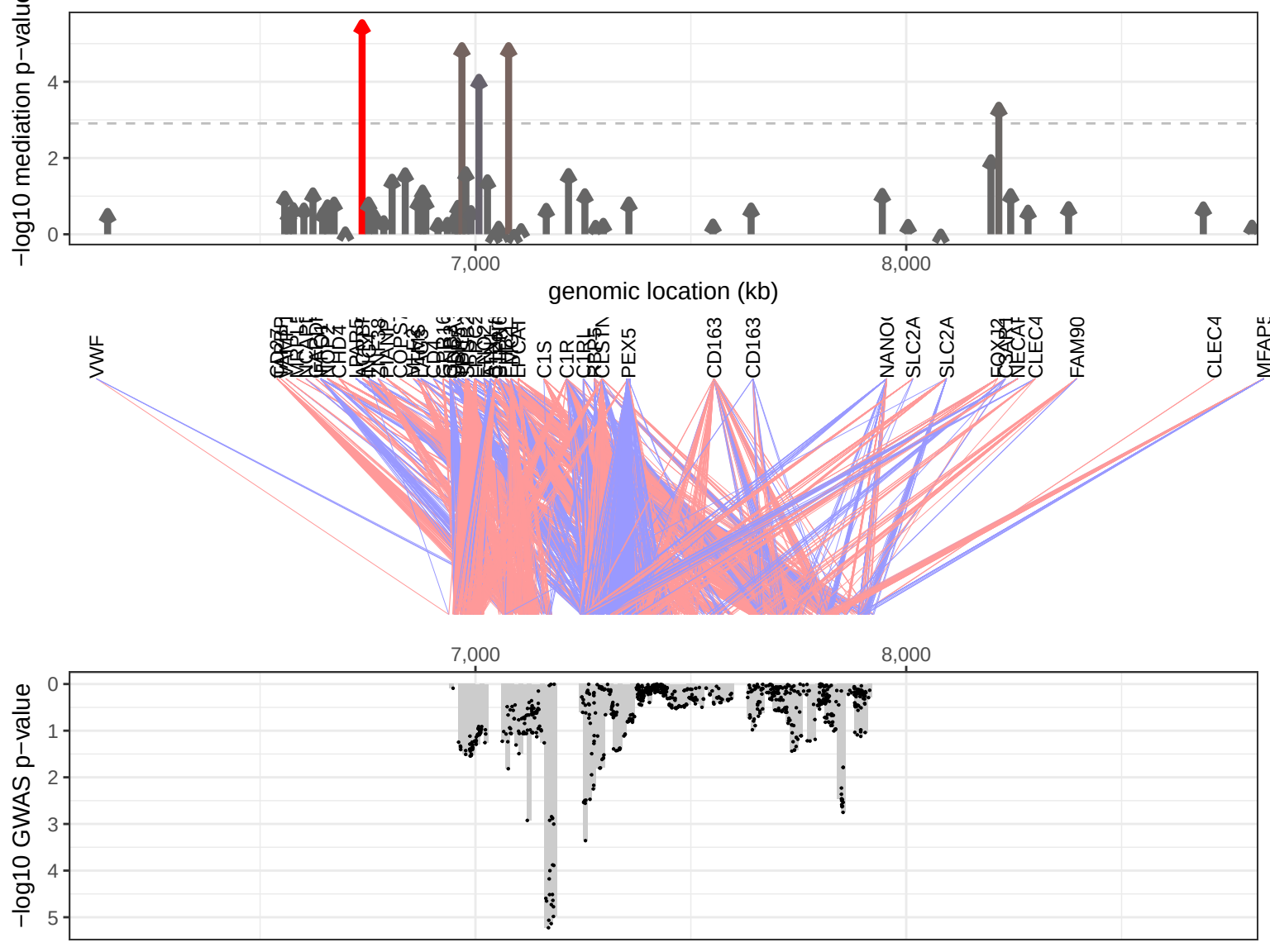
−log10 mediation p−value
0
2
4
7,000
8,000
genomic location (kb)
VWF
C1S
C1R
PEX5
CD163
CD163
SLC2A
SLC2A
CLEC4
FAM90
CLEC4
7,000
8,000
−log10 GWAS p−value
0
1
2
3
4
5