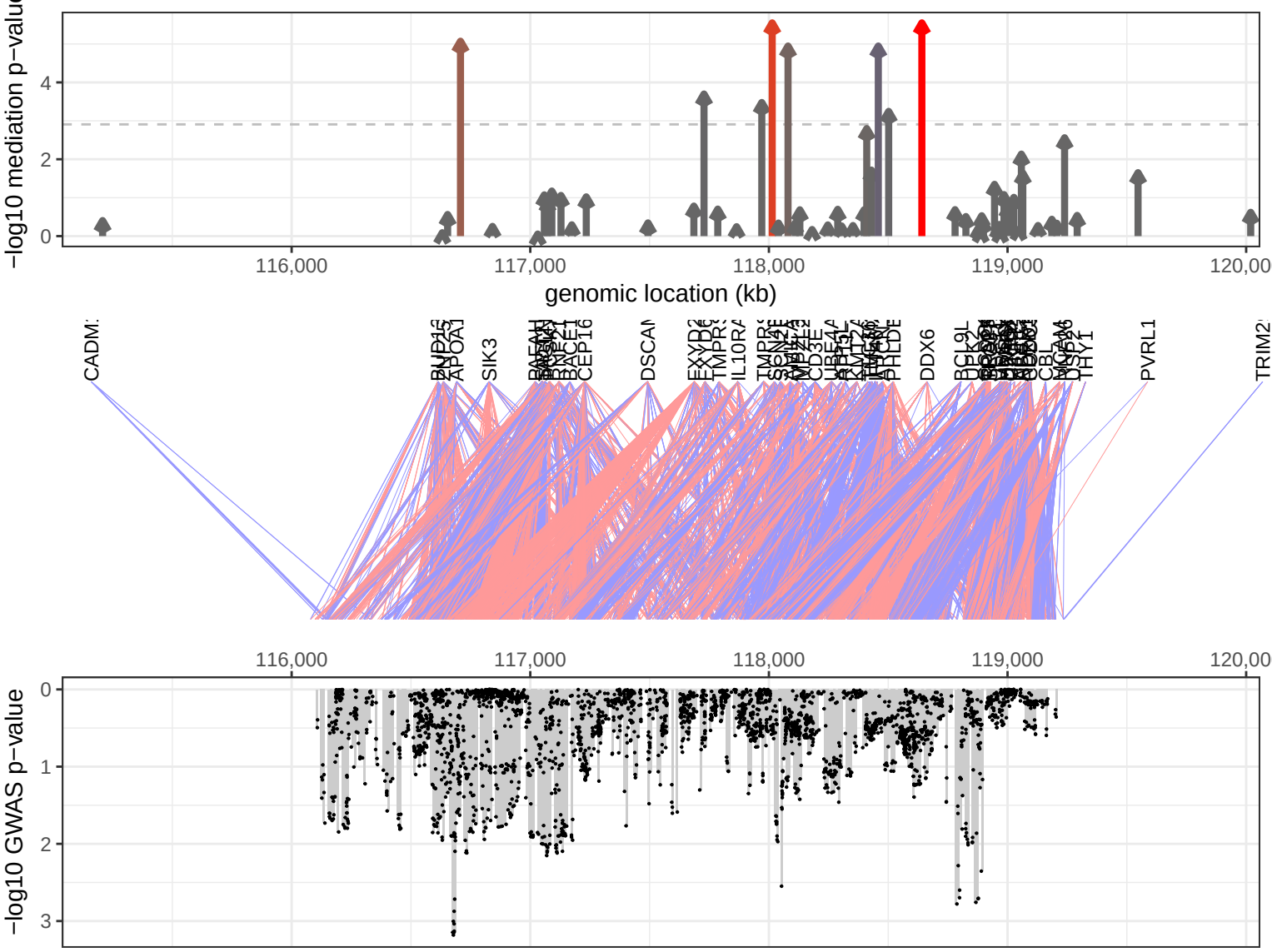
−log10 mediation p−value
0
2
4
116,000
117,000
118,000
119,000
120,00
genomic location (kb)
CADM
SIK3
DSCAM
IL10RA
DDX6
PVRL1
TRIM2
116,000
117,000
118,000
119,000
120,00
−log10 GWAS p−value
0
1
2
3