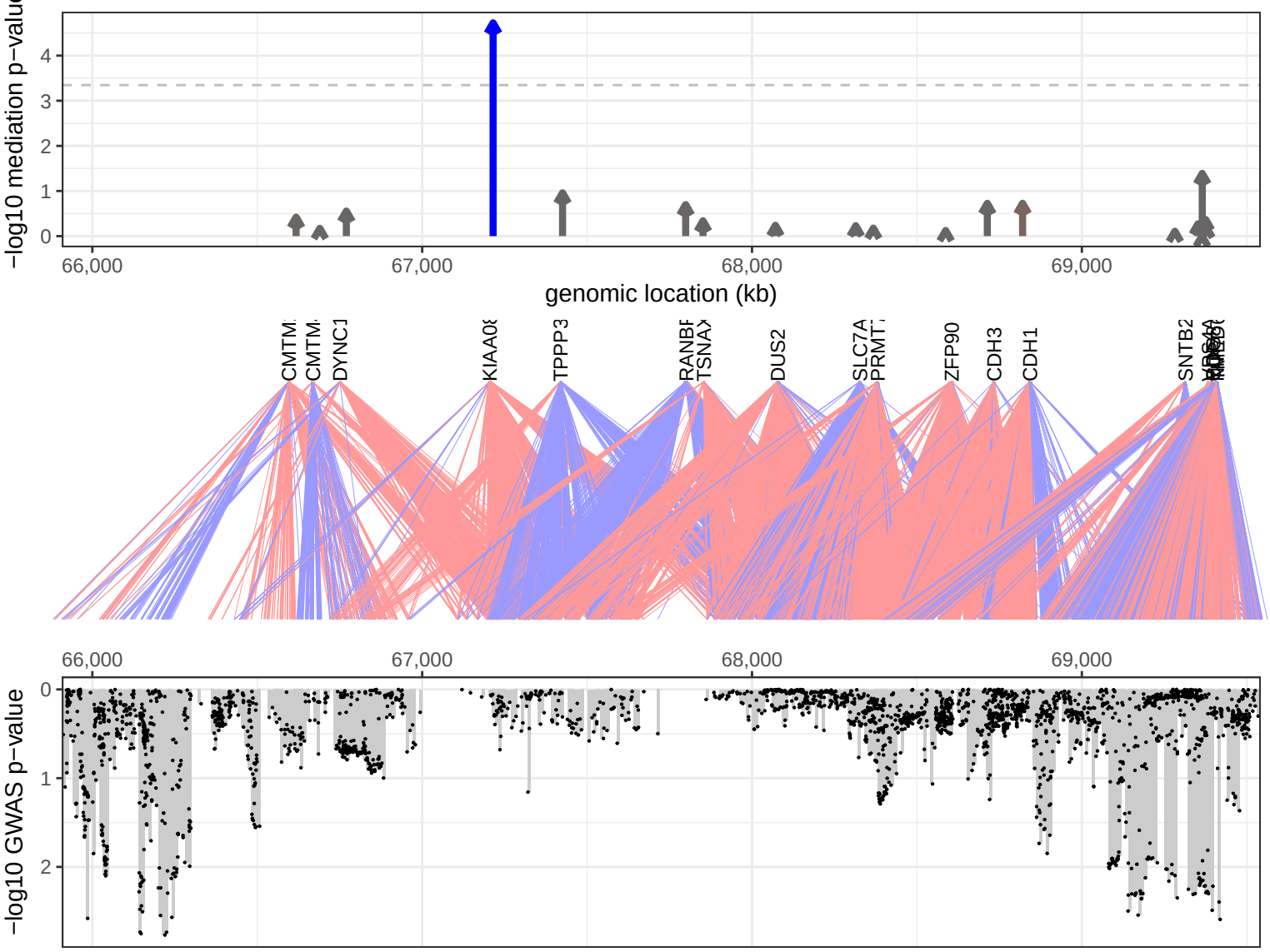
−log10 mediation p−value
4
3
2
1
0
66,000
67,000
68,000
69,000
genomic location (kb)
CMTM
CMTM
DYNC1
KIAA0
TPPP3
RANBP
TSNAX
DUS2
SLC7A
PRMT
ZFP90
CDH3
CDH1
SNTB2
−log10 GWAS p−value
0
1
2