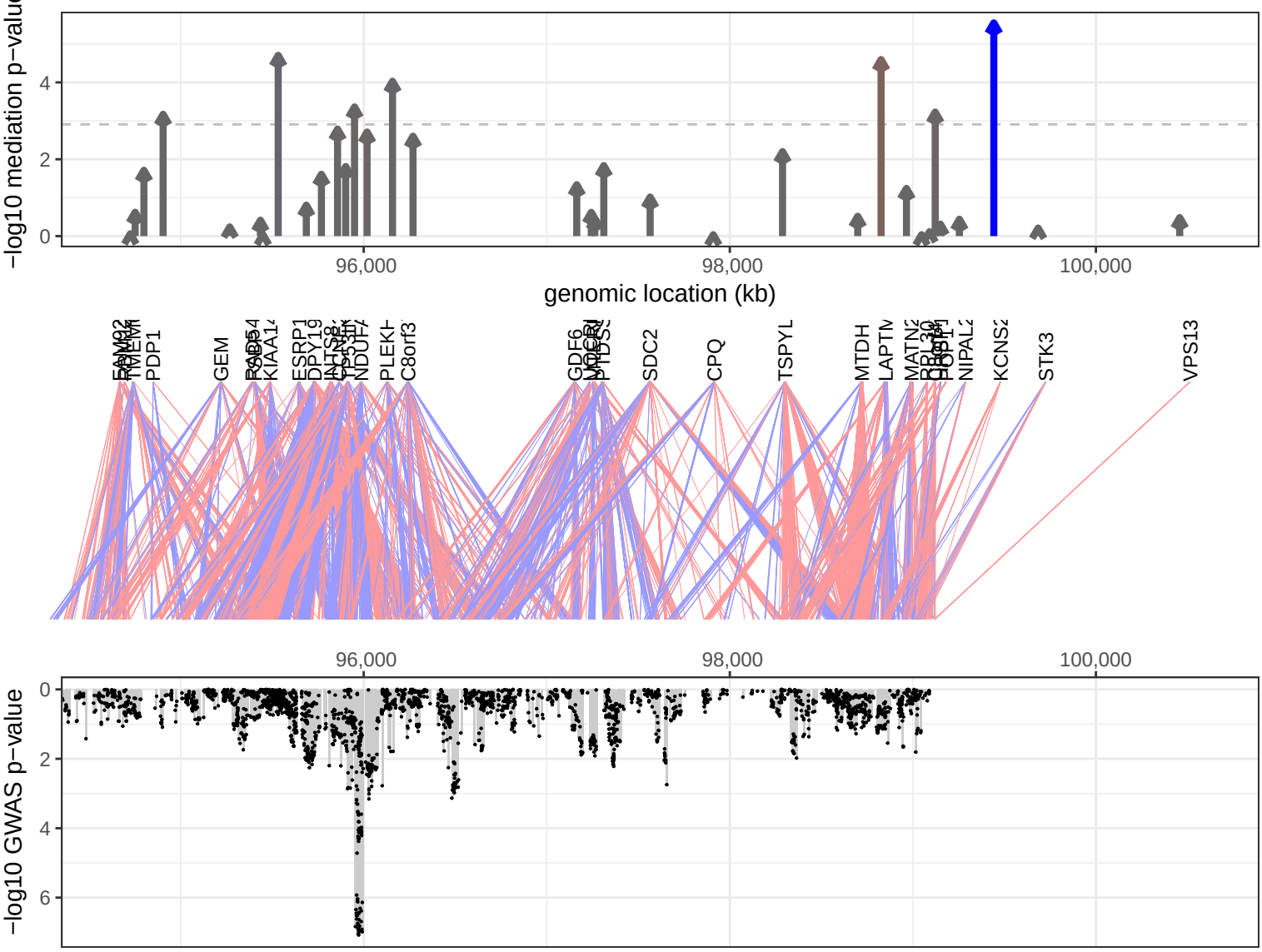
−log10 mediation p−value
0
2
4
96,000
98,000
100,000
genomic location (kb)
PDP1
GEM
KIAA1
ESRP1
DPY19
NDUFA
PLEKH
C8orf3
GDF6
SDC2
CPQ
TSPYL
MTDH
LAPTM
NIPAL
KCNS
STK3
VPS13
96,000
98,000
100,000
−log10 GWAS p−value
0
2
4
6