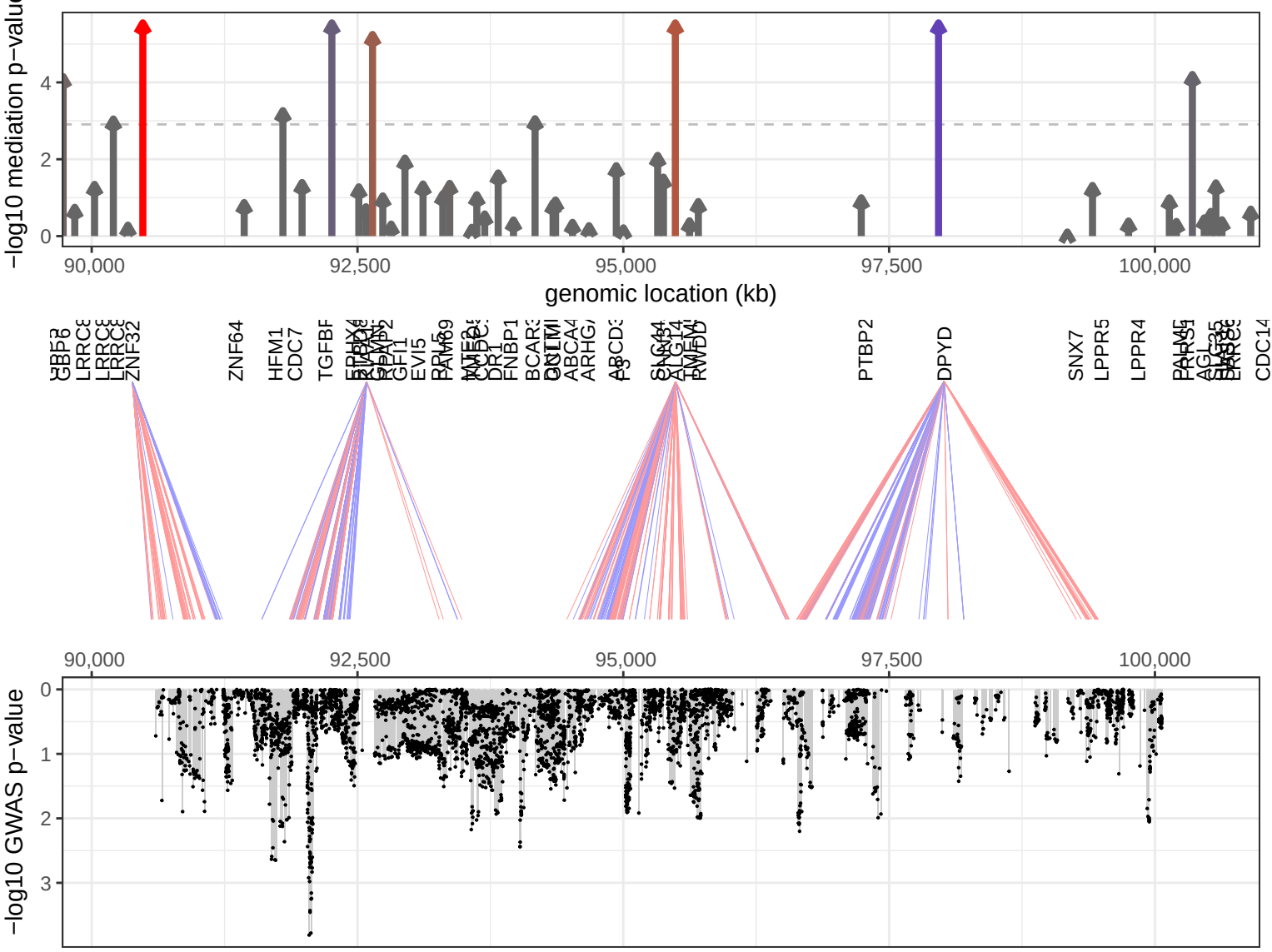
−log10 mediation p−value
0
2
4
90,000
92,500
95,000
97,500
100,000
genomic location (kb)
ZNF32
ZNF64
HFM1
CDC7
TGFBF
EVI5
FNBP1
PTBP2
DPYD
SNX7
LPPR5
LPPR4
−log10 GWAS p−value
0
1
2
3