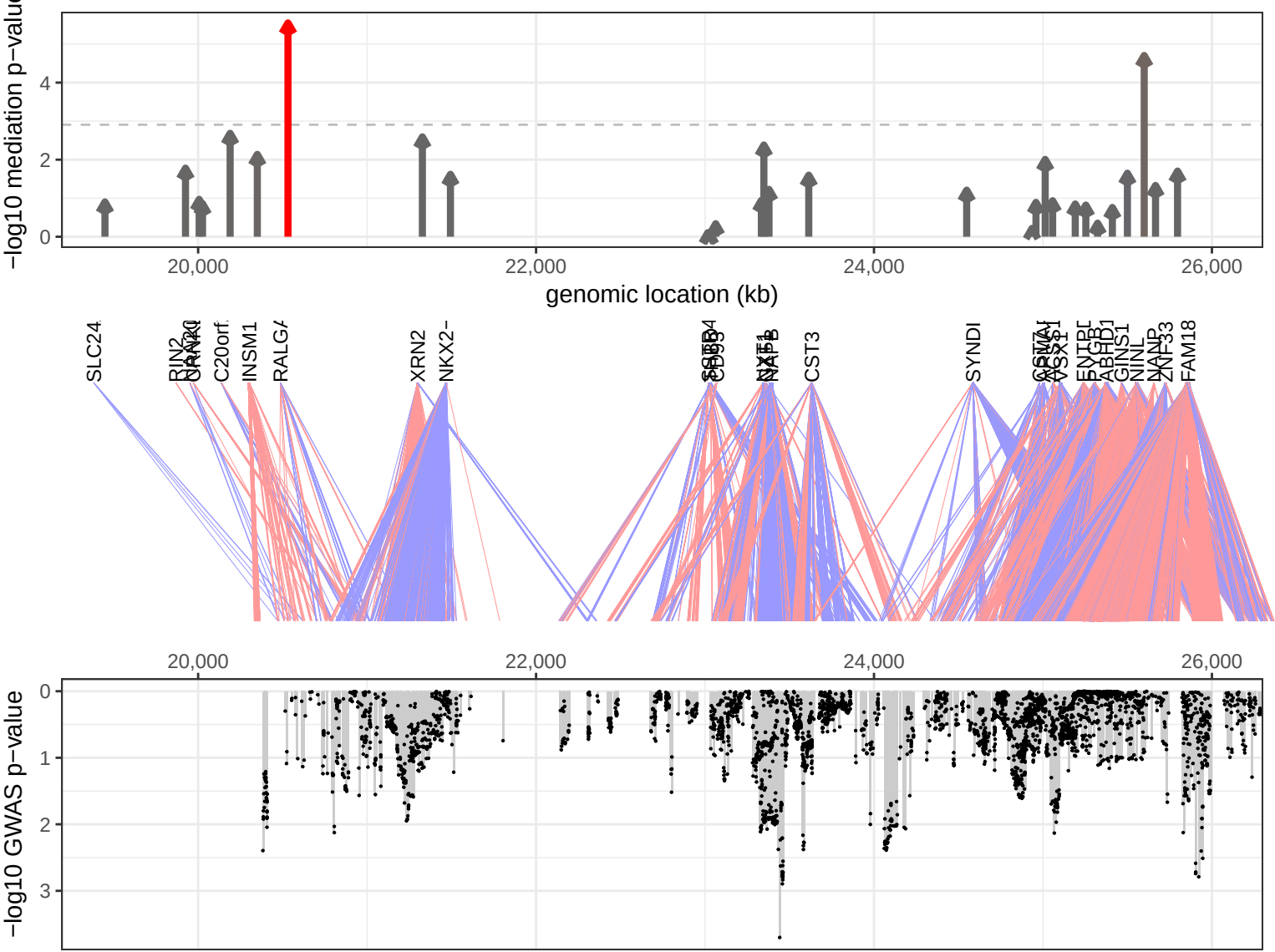
−log10 mediation p−value
0
2
4
20,000
22,000
24,000
26,000
genomic location (kb)
SLC24
INSM1
XRN2
NKX2−
CST3
SYNDI
FAM18
20,000
22,000
24,000
26,000
−log10 GWAS p−value
0
1
2
3