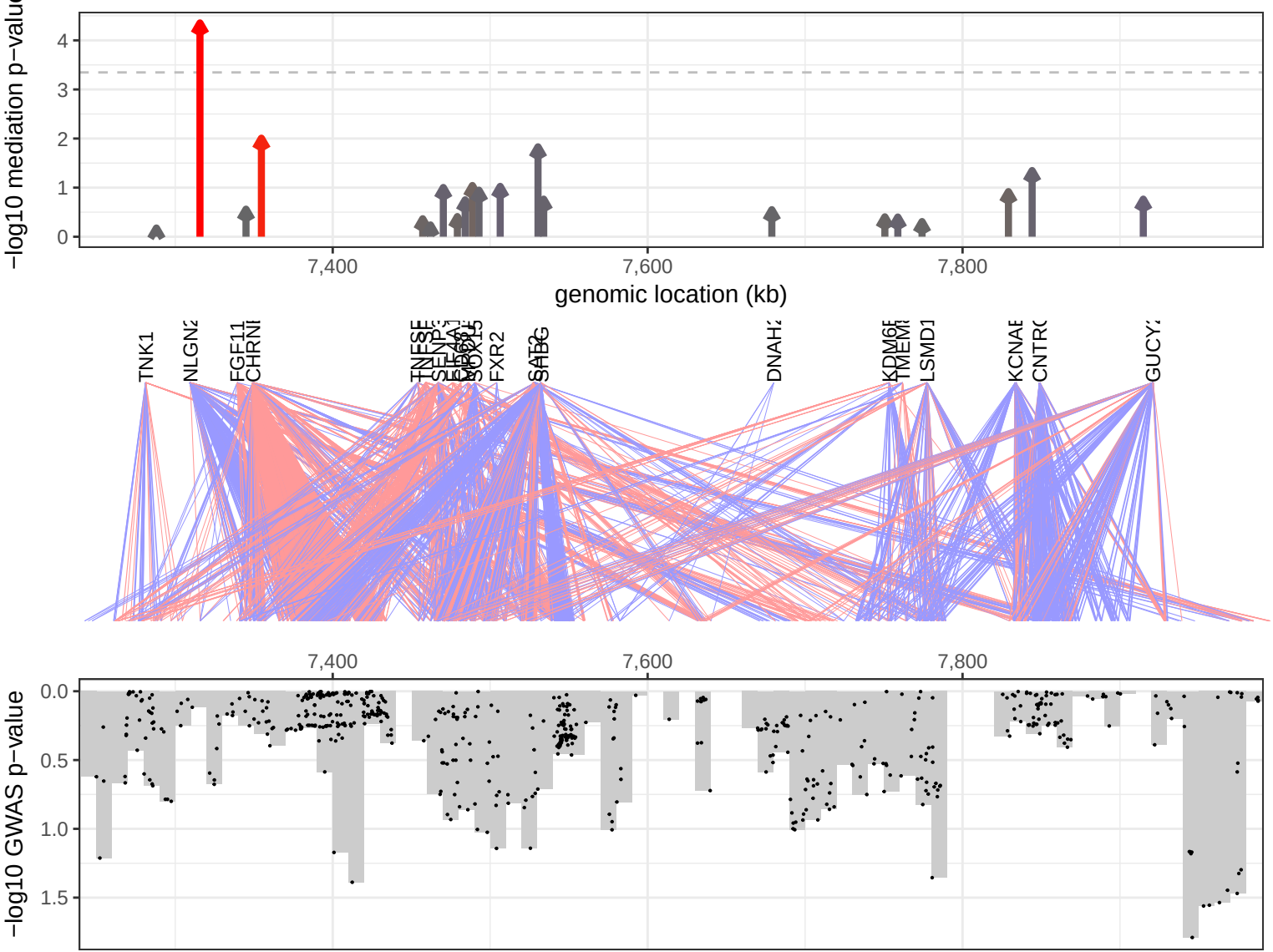
−log10 mediation p−value
4
3
2
1
0
7,400
7,600
7,800
genomic location (kb)
TNK1
NLGN2
FGF11
CHRNI
FXR2
DNAH2
LSMD1
KCNAE
CNTRC
GUCY2
7,400
7,600
7,800
−log10 GWAS p−value
0.0
0.5
1.0
1.5