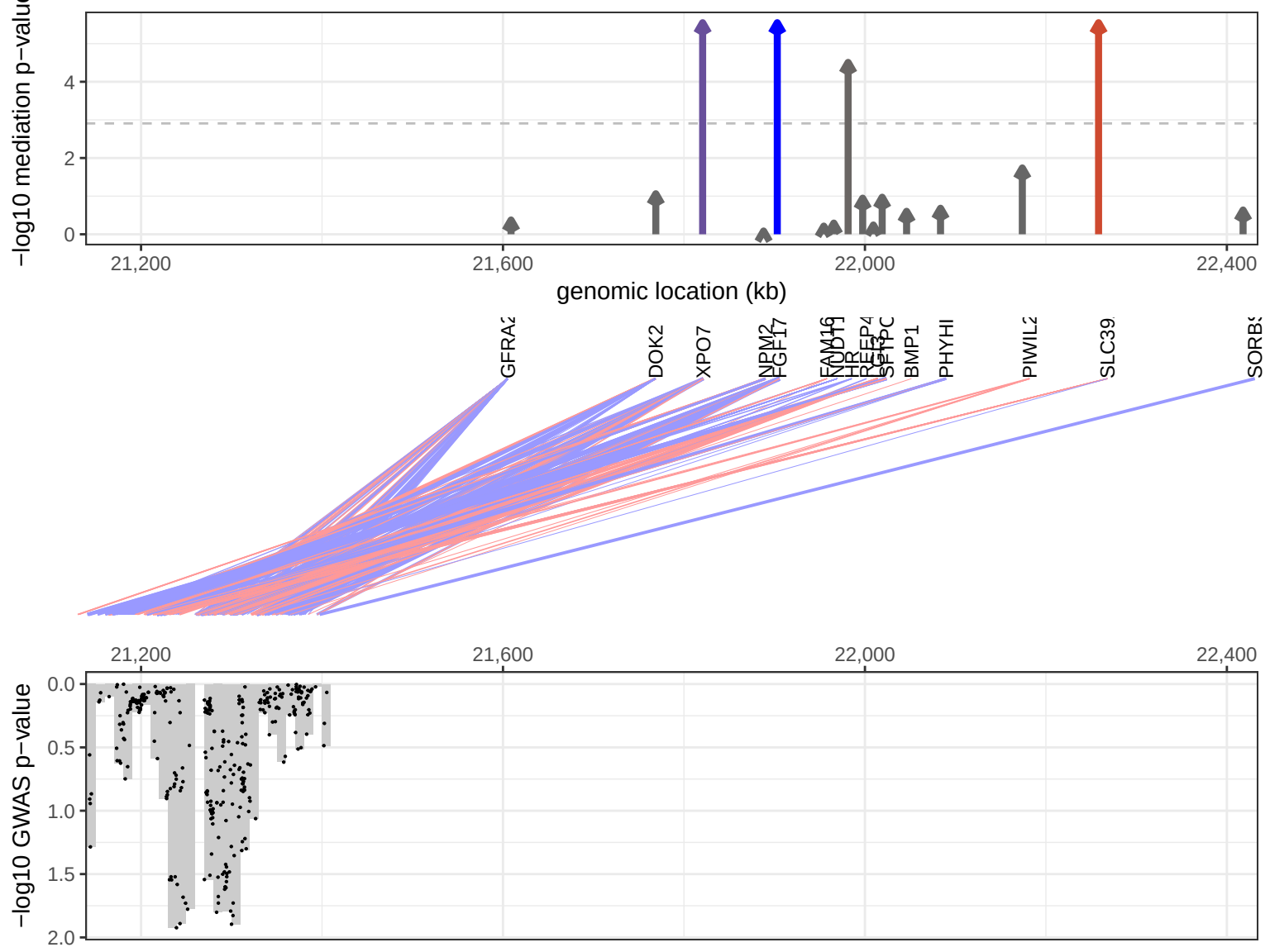
−log10 mediation p−value
0
2
4
21,200
21,600
22,000
22,400
genomic location (kb)
GFRA2
DOK2
XPO7
NPM2
FGF17
FAM46
NUDT11
HR
RGS5
REEP4
SFTPC
BMP1
PHYHI
PIWIL2
SLC39
SORBS
21,200
21,600
22,000
22,400
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0