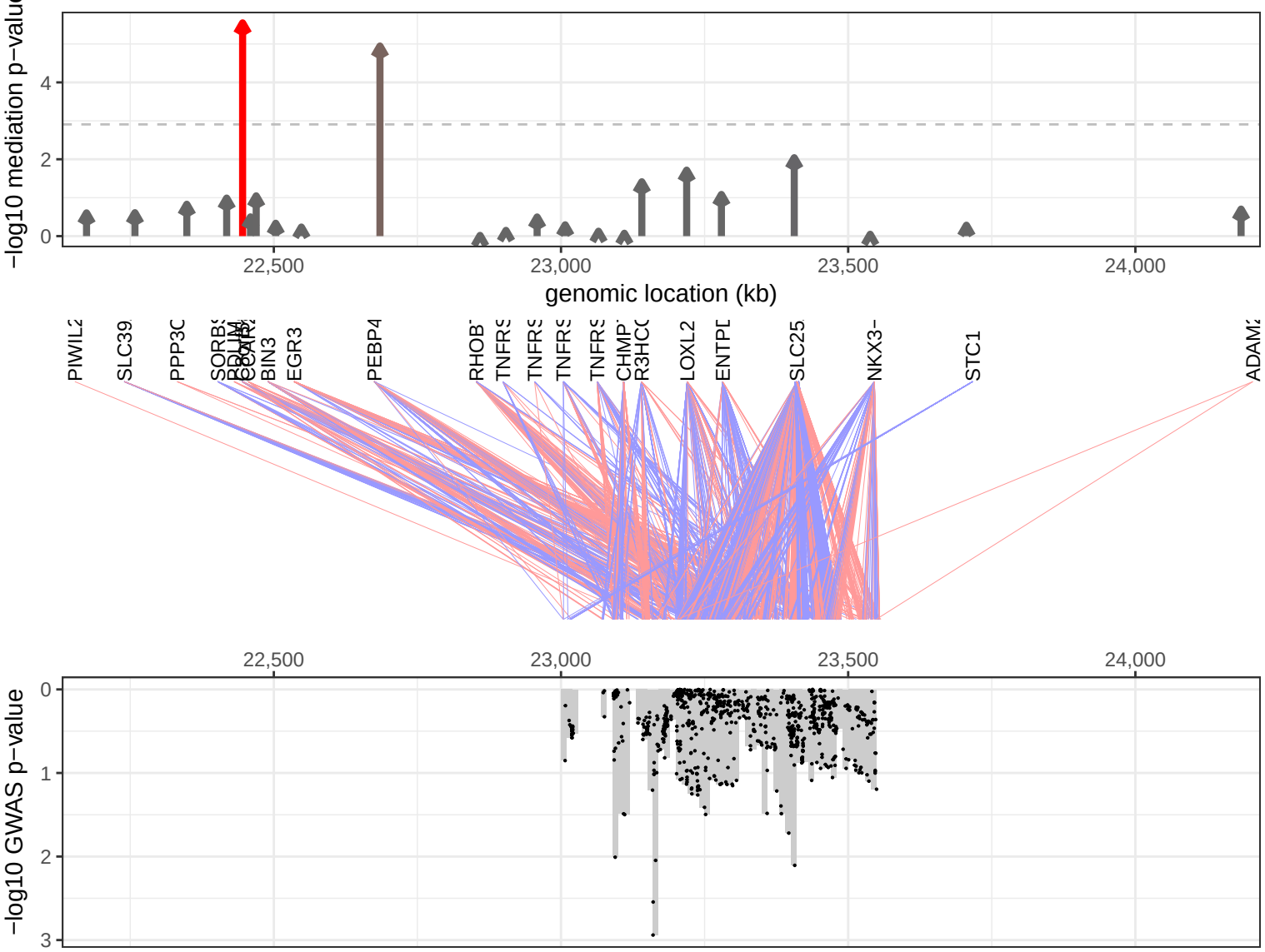
−log10 mediation p−value
0
2
4
22,500
23,000
23,500
24,000
genomic location (kb)
PIWIL2
SLC39
PPP3C
BIN3
EGR3
PEBP4
LOXL2
SLC25
NKX3-
STC1
ADAM
22,500
23,000
23,500
24,000
−log10 GWAS p−value
0
1
2
3