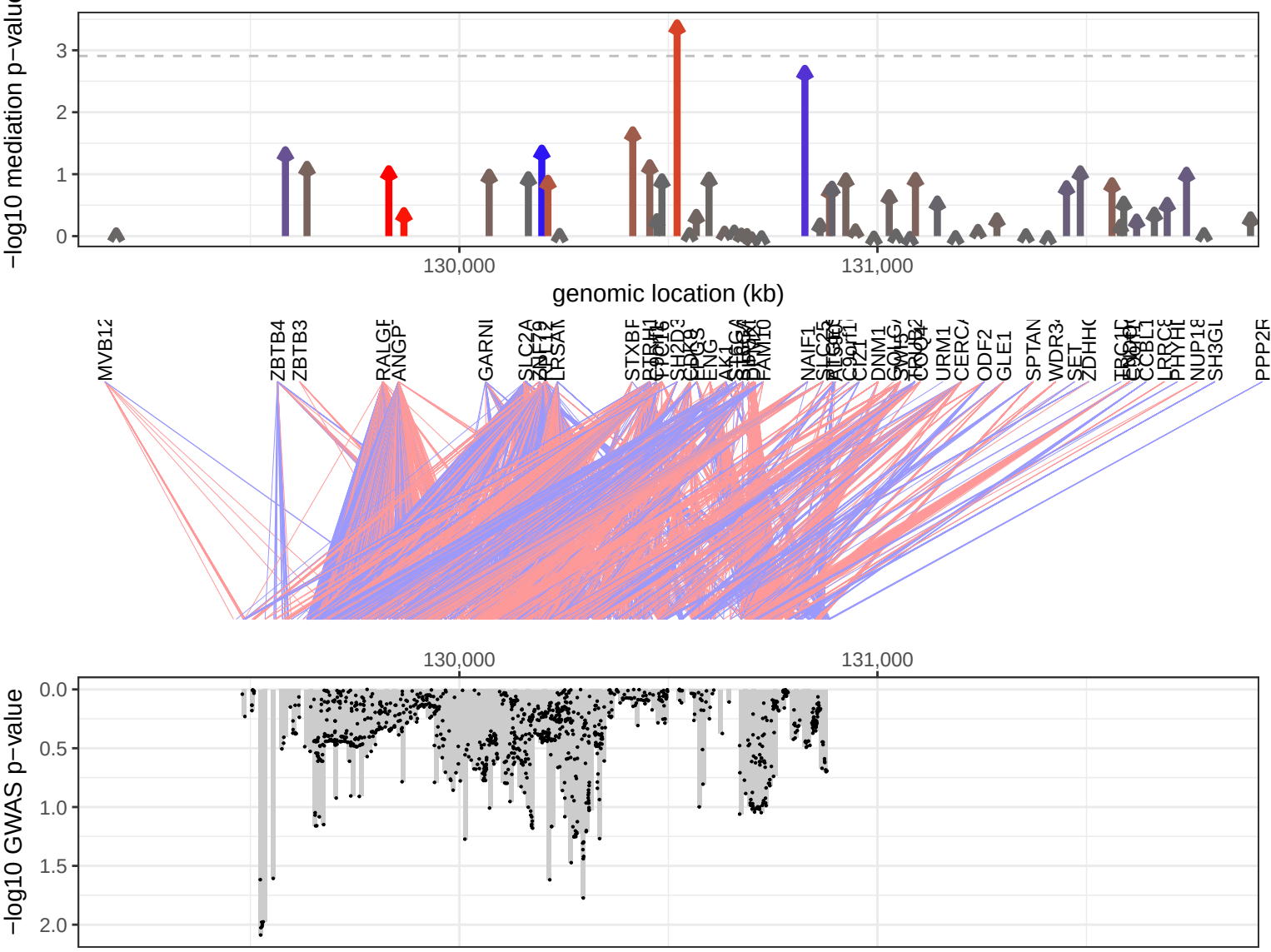
−log10 mediation p−value
3
2
1
0
130,000
131,000
genomic location (kb)
ZBTB4
ZBTB3
RALGF
ANGP
GARNI
STXBF
ENG
NAIF1
DNM1
URM1
ODF2
GLE1
SPTAN
WDR3
SET
NUP18
SH3GL
PPP2R
130,000
131,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0