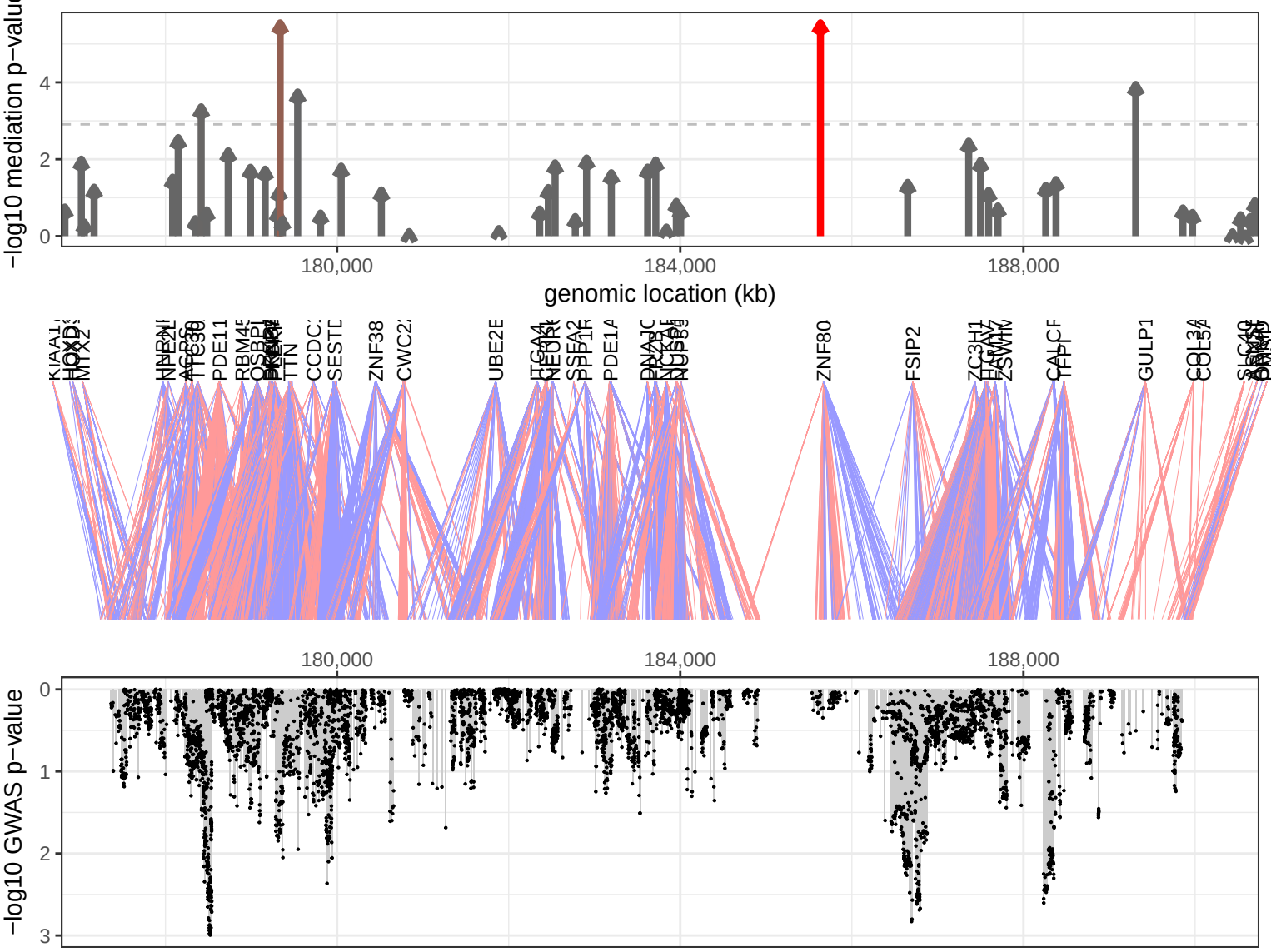
−log10 mediation p−value
0
2
4
180,000
184,000
188,000
genomic location (kb)
KIAA1
HOXD
MTX2
NPEPL
ACP6
TC30
PDE11
RBM45
OSBP
TTN
CCDC
SESTD
ZNF38
CWC27
UBE2E
ITGA1
NEURL
SSFA2
PPP1R
PDE1A
PPAT
NCAM
NUP
ZNF80
FSIP2
ZC3H1
ZSWIM
CALCF
TFPT
GULP1
COL3A
−log10 GWAS p−value
0
1
2
3