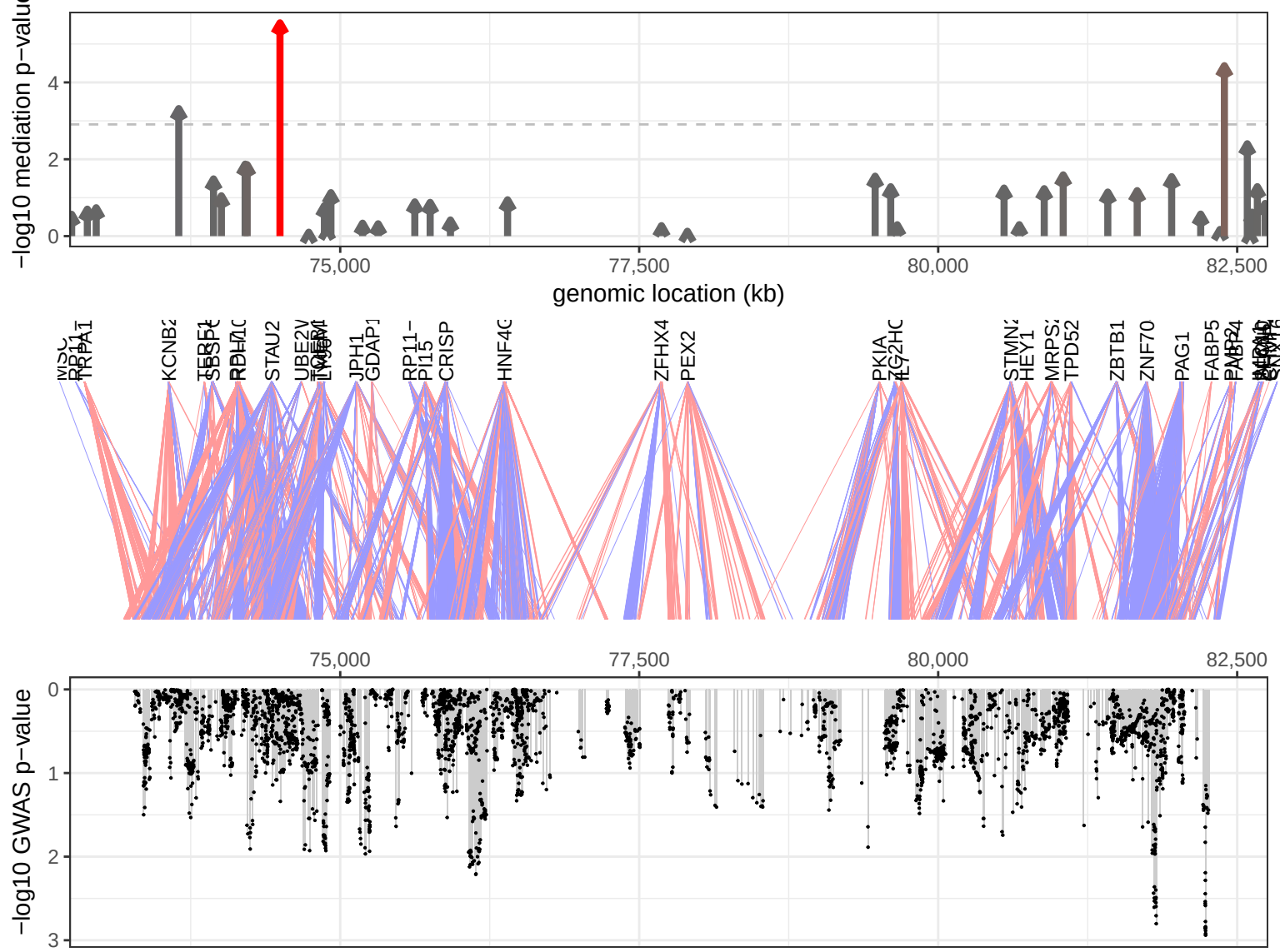
−log10 mediation p−value
0
2
4
75,000
77,500
80,000
82,500
genomic location (kb)
KCNB2
STAU2
JPH1
GDAP1
RP11−
PI15
CRISP
HNF4G
ZFHX4
PEX2
PKIA
STMN2
HEY1
MRPS2
TPD52
ZBTB1
ZNF70
PAG1
FABP5
−log10 GWAS p−value
0
1
2
3