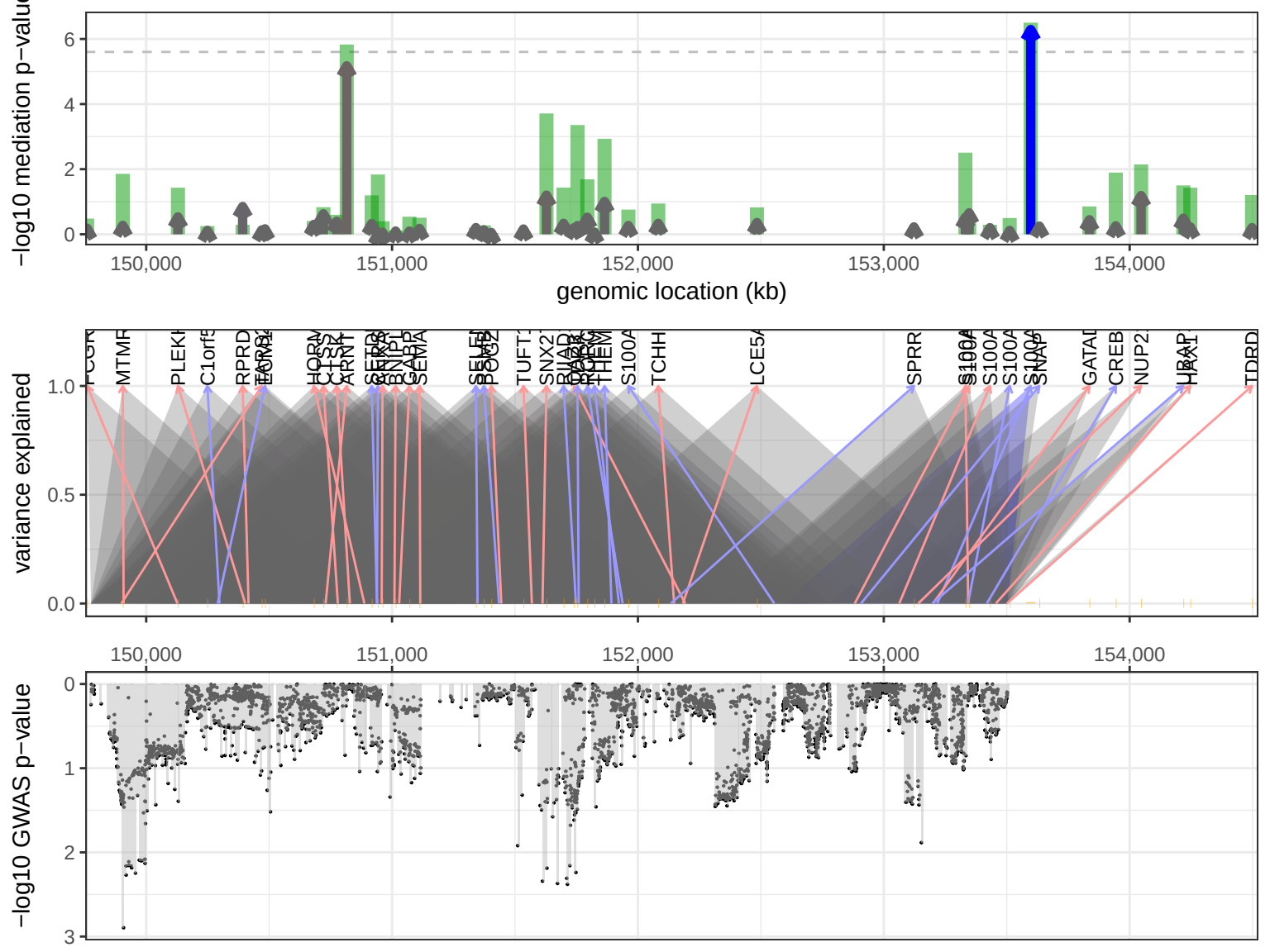
−log10 mediation p−value
genomic location (kb)
variance explained
−log10 GWAS p−value
150,000
151,000
152,000
153,000
154,000
MTMR
PLEKH
C1orf5
RPRD
TUFT1
SNX2
S100A
TCHH
LCE5A
SPRR
S100A
S100A
S100A
GATAD
CREB
NUP2
TDRD