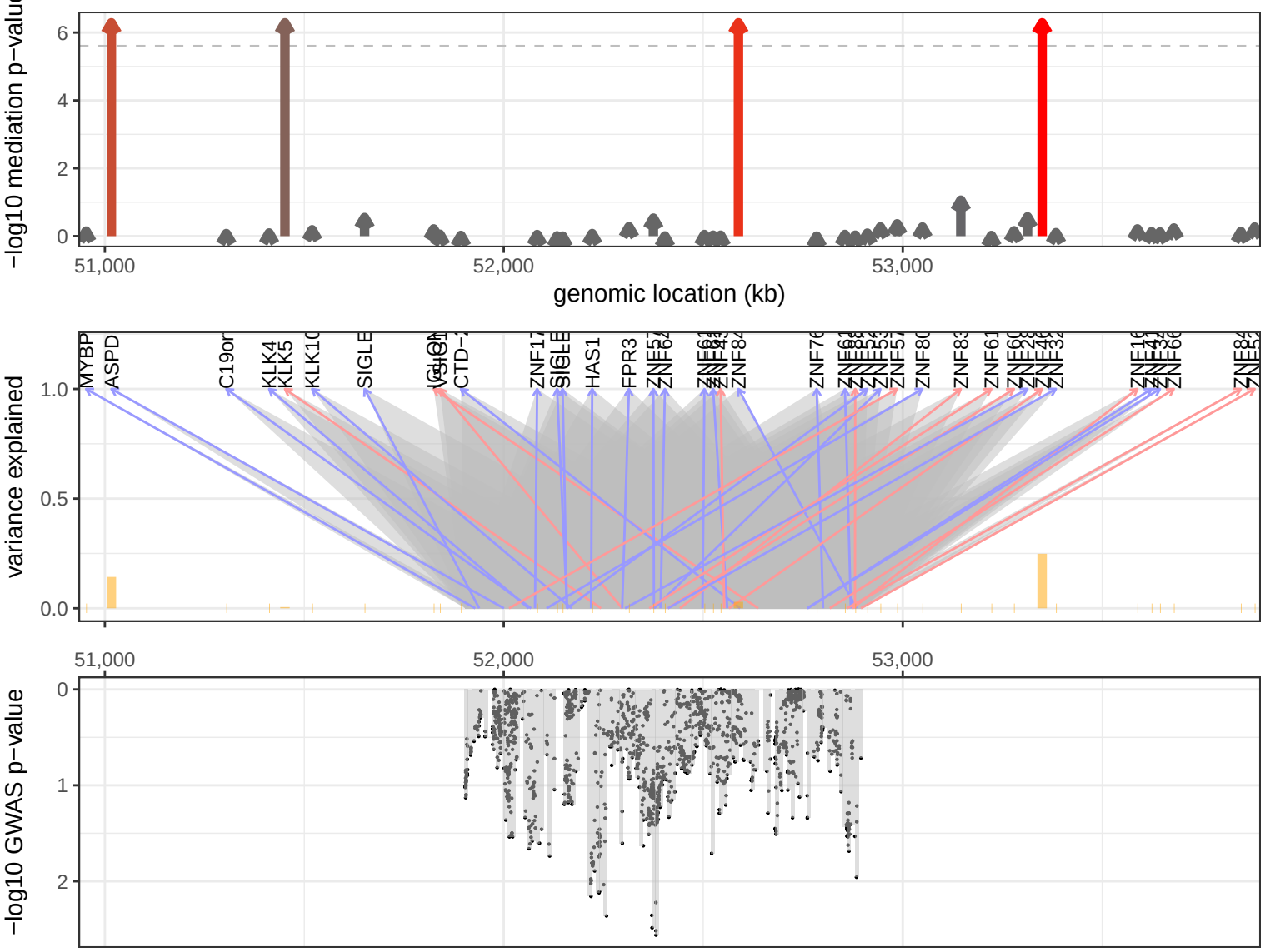
−log10 mediation p−value
6
4
2
0
51,000
52,000
53,000
genomic location (kb)
variance explained
1.0
0.5
0.0
MYBP
ASPD
C19orf
KLK4
KLK5
SIGLEC
CTD-
ZNF17
HAS1
FPR3
ZNF84
ZNF76
ZNF57
ZNF80
ZNF83
ZNF61
ZNF32
−log10 GWAS p−value
0
1
2