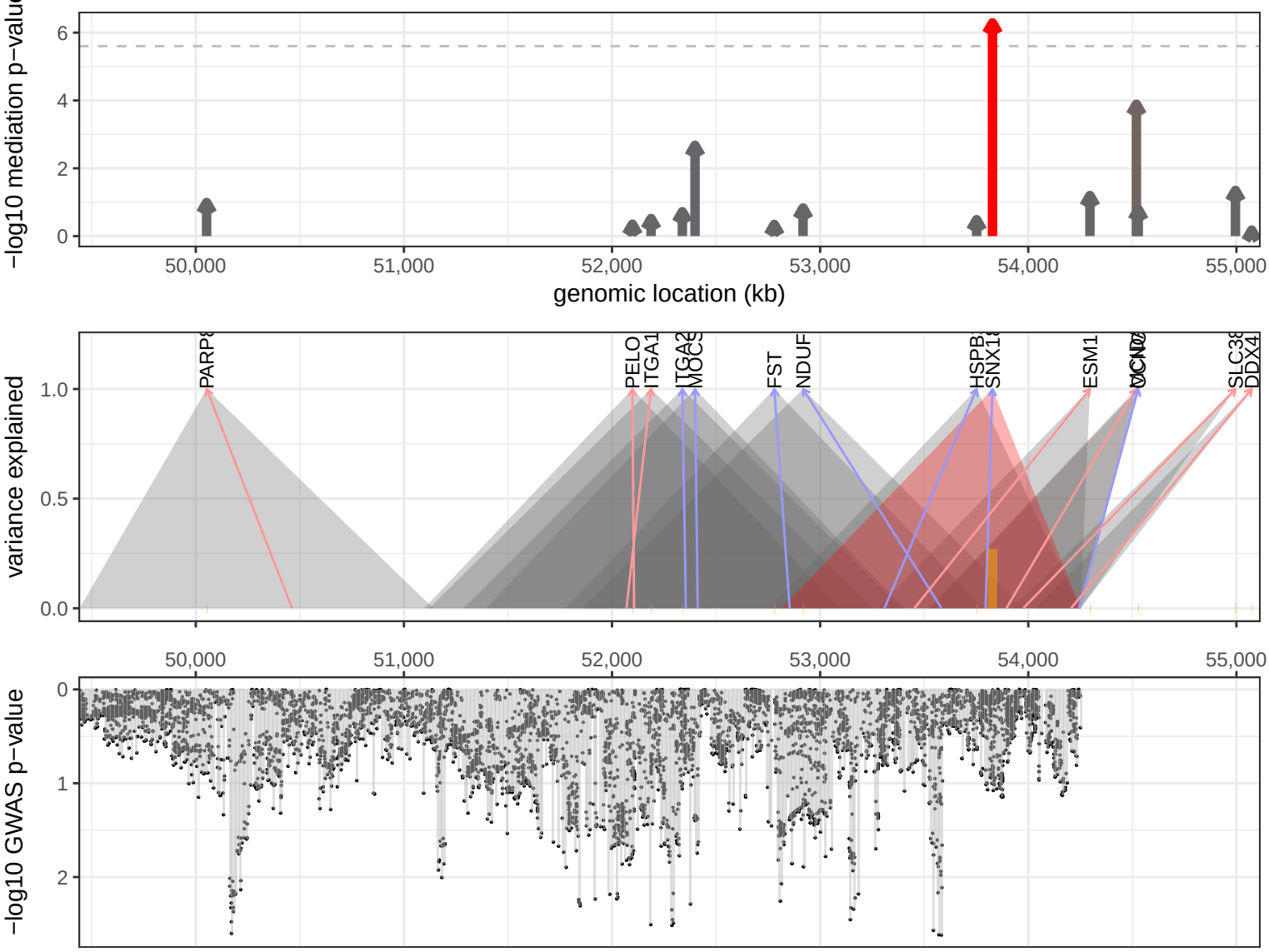
−log10 mediation p−value
6
4
2
0
50,000
51,000
52,000
53,000
54,000
55,000
genomic location (kb)
variance explained
1.0
0.5
0.0
PARP8
PELO
ITGA1
ITGA2
MOCS
FST
NDUF
HSPB
SNX1
ESM1
SLC38
DDX4
−log10 GWAS p−value
0
1
2