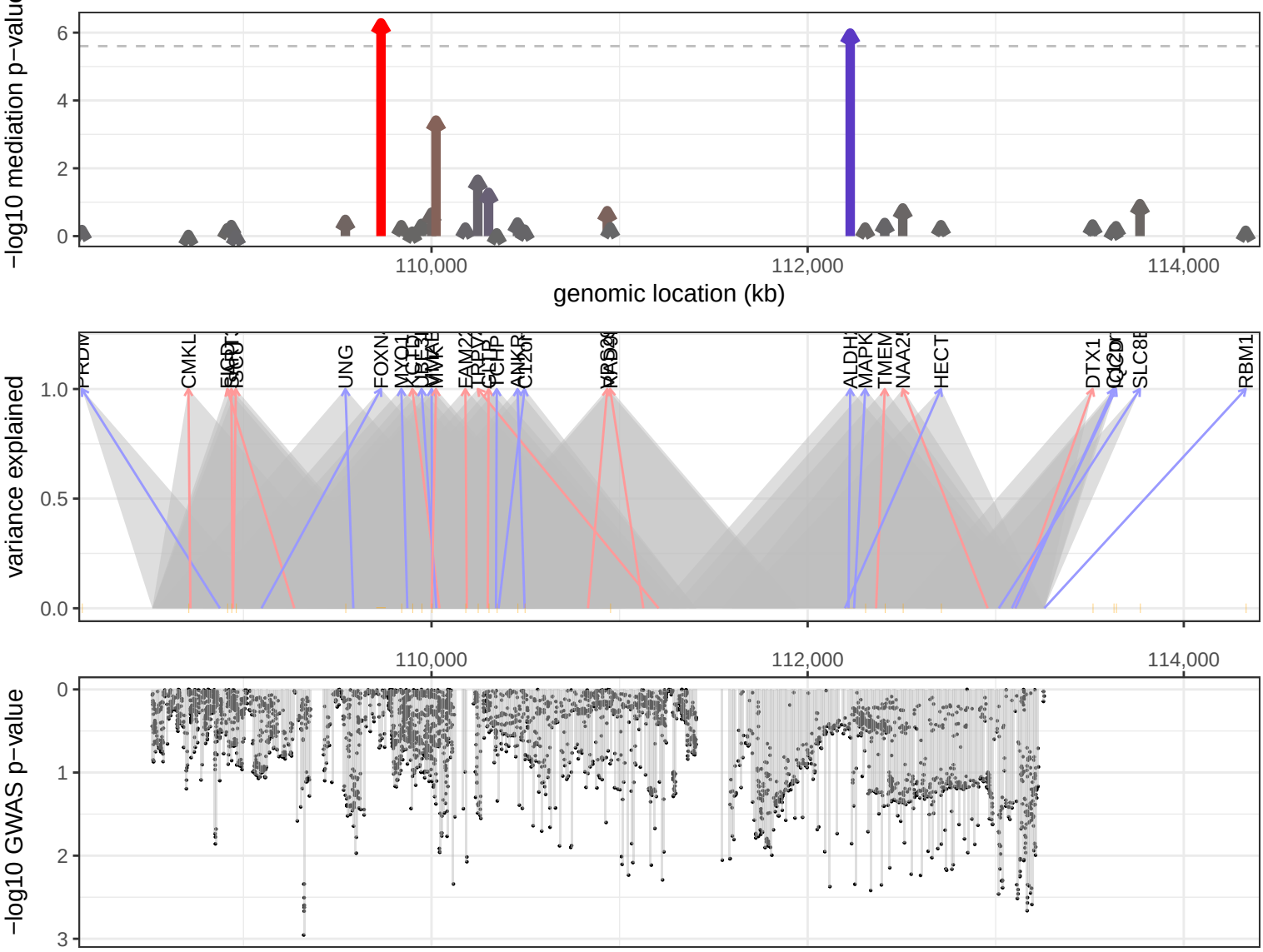
−log10 mediation p−value
6
4
2
0
110,000
112,000
114,000
genomic location (kb)
variance explained
1.0
0.5
0.0
CMKL
UNG
FOXN
ALDH
MAPK
TMEM
NAA2
HECT
DTX1
SLC8
RBM1
110,000
112,000
114,000
−log10 GWAS p−value
0
1
2
3