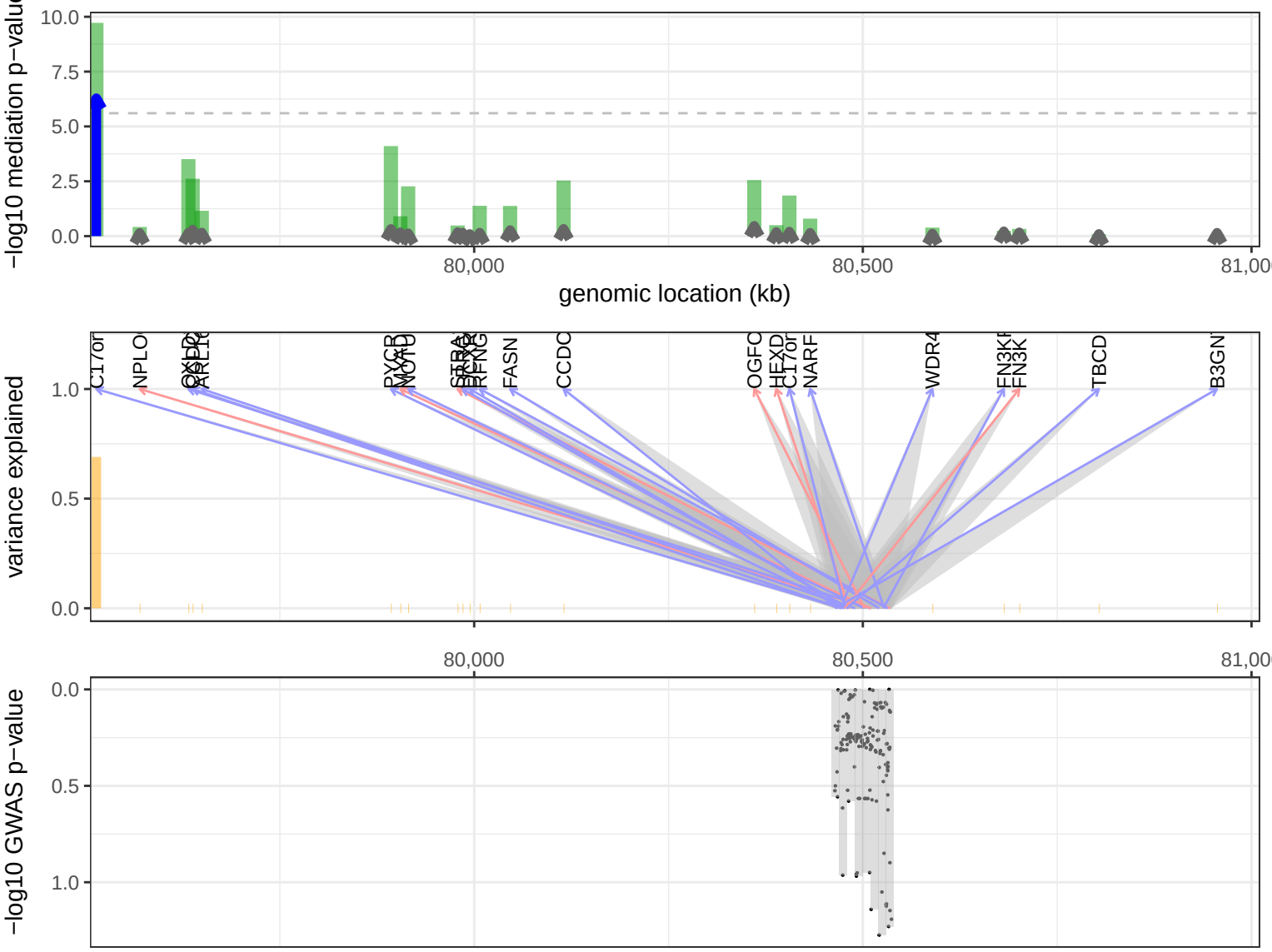
−log10 mediation p−value
10.0
7.5
5.0
2.5
0.0
80,000
80,500
genomic location (kb)
variance explained
1.0
0.5
0.0
NPLO
FASN
CCDC
NARF
WDR4
TBCD
B3GN
−log10 GWAS p−value
0.0
0.5
1.0