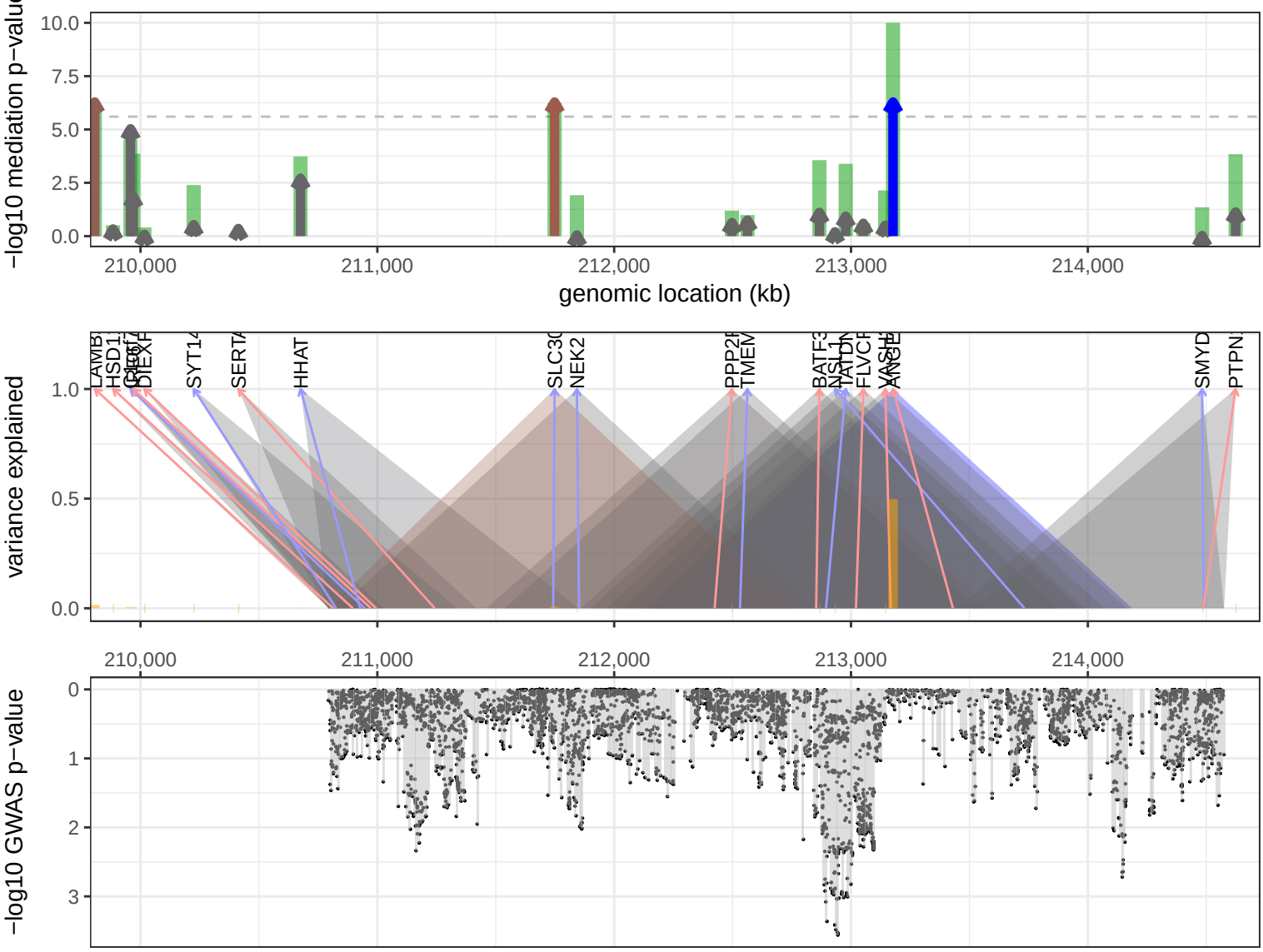
−log10 mediation p−value
10.0
7.5
5.0
2.5
0.0
210,000
211,000
212,000
213,000
214,000
genomic location (kb)
variance explained
1.0
0.5
0.0
SYT14
SERTA
HHAT
SLC30
NEK2
PPP2R
TMEM
BATF3
TATDN
FLVCR
SMYD
PTPN
−log10 GWAS p−value
0
1
2
3