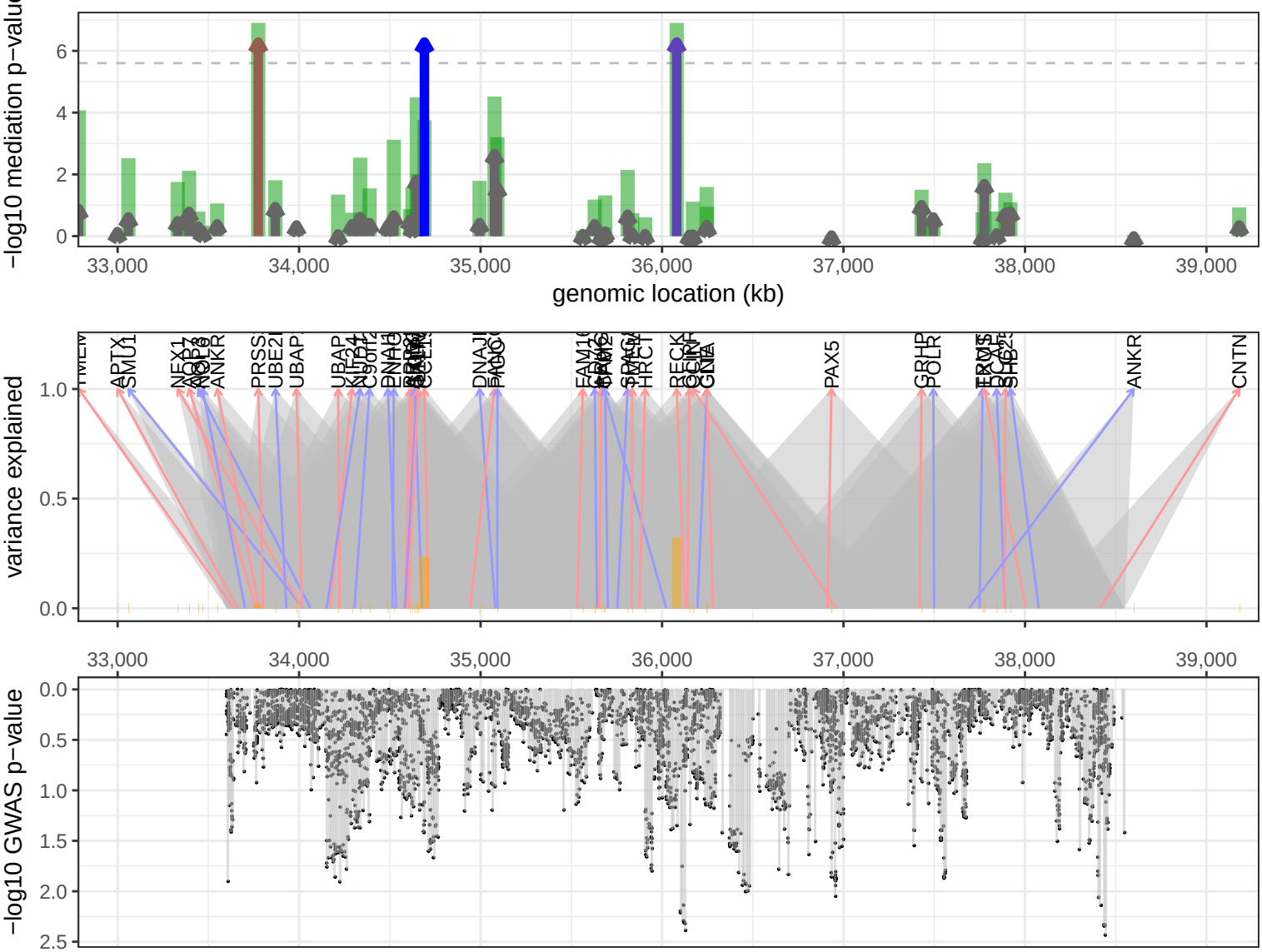
−log10 mediation p−value
genomic location (kb)
variance explained
−log10 GWAS p−value
33,000
34,000
35,000
36,000
37,000
38,000
39,000
PAX5
ANKR
CNTN