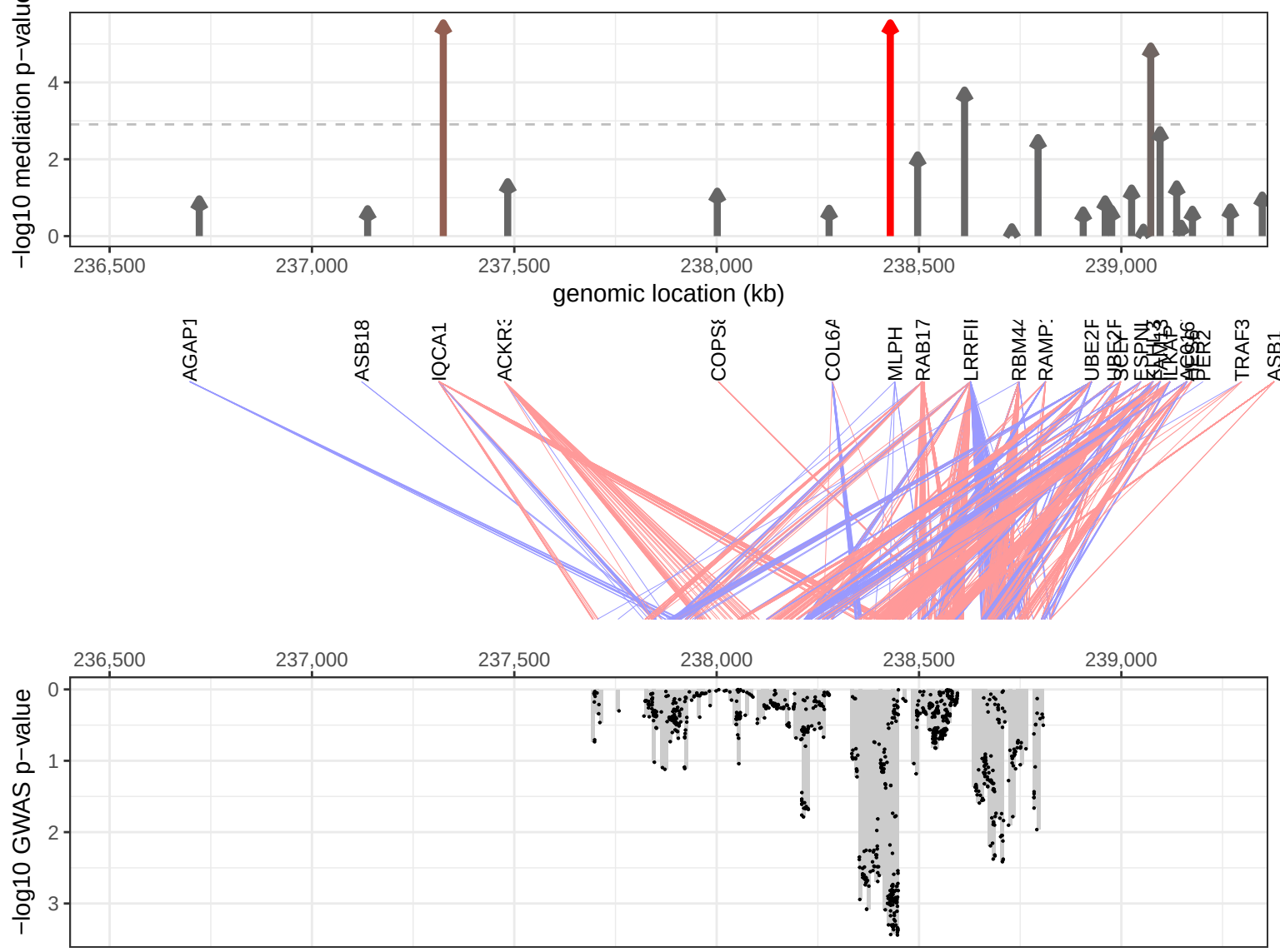
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2
4
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237,000
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genomic location (kb)
AGAP1
ASB18
IQCA1
ACKR3
COPS8
COL6A
MLPH
RAB17
LRRFIP
RBM42
RAMP1
UBE2F
UBE2F
ESPN
TRAF3
ASB1
236,500
237,000
237,500
238,000
238,500
239,000
−log10 GWAS p−value
0
1
2
3