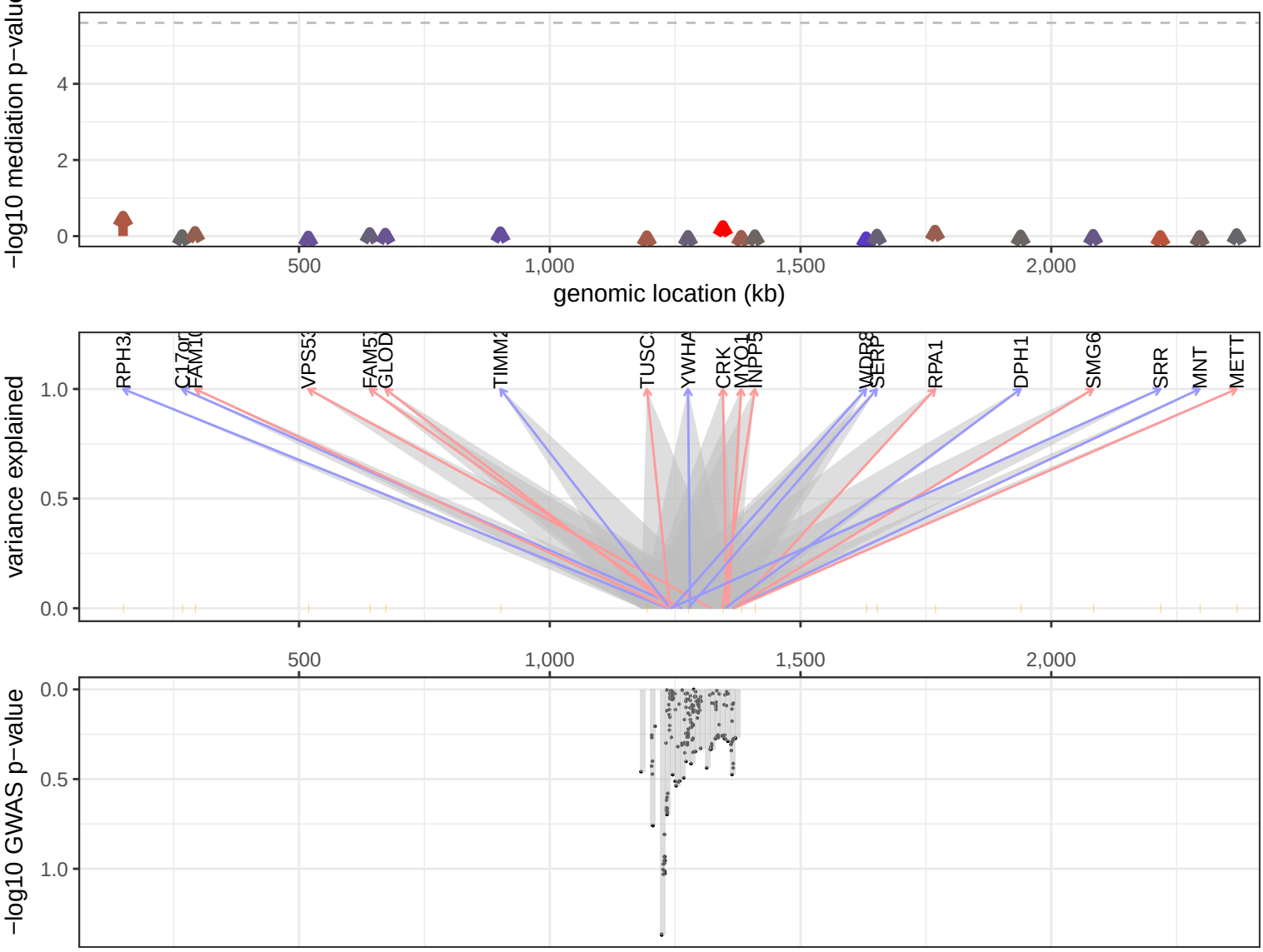
−log10 mediation p−value
4
2
0
500
1,000
1,500
2,000
genomic location (kb)
variance explained
1.0
0.5
0.0
RPH3
VPS5
FAM5
GLOD
TIMM2
TUSC
YWHA
CRK
MYO1
NPP5
SERP
RPA1
DPH1
SMG6
SRR
MNT
METT
500
1,000
1,500
2,000
−log10 GWAS p−value
0.0
0.5
1.0