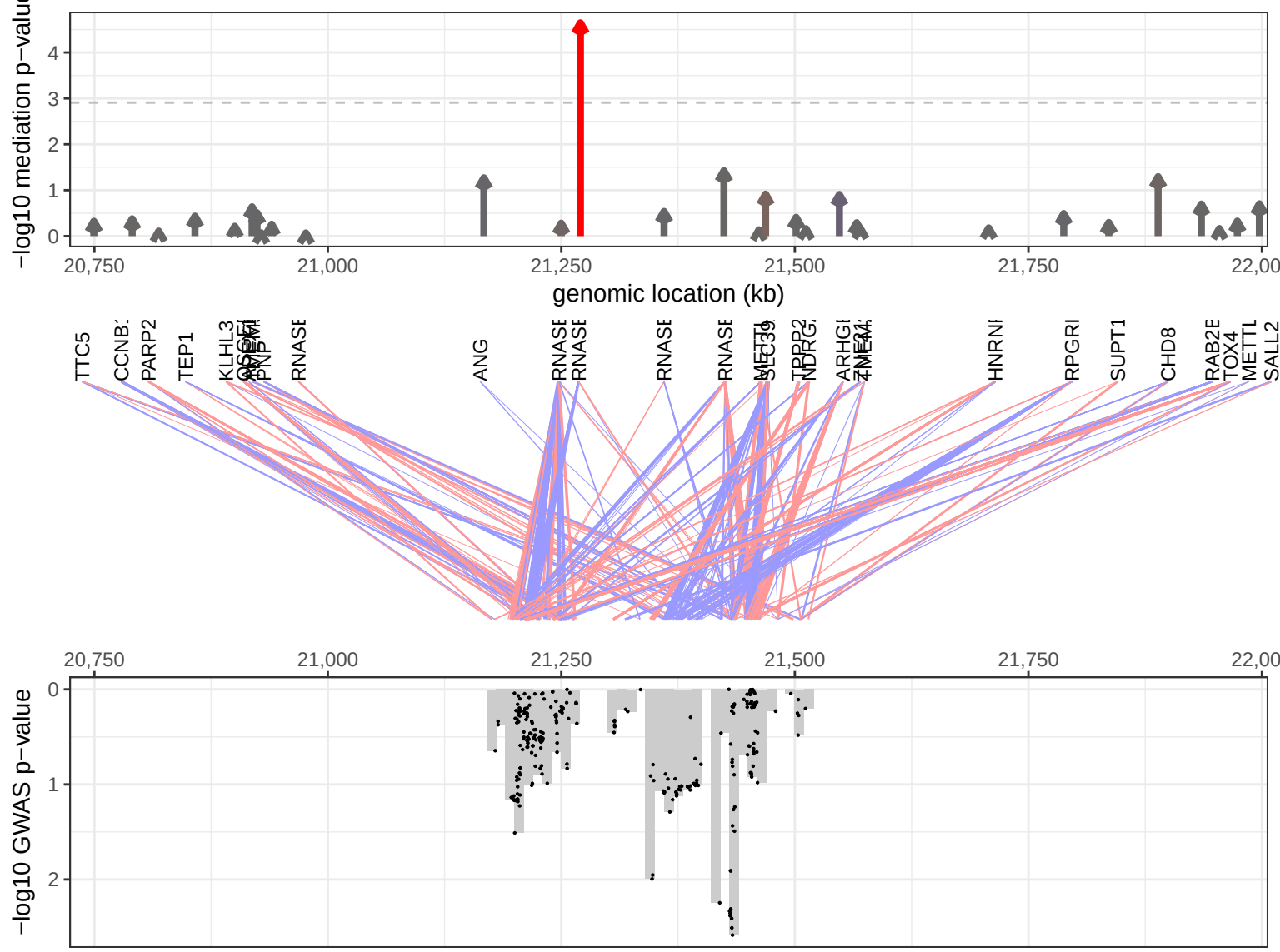
−log10 mediation p−value
genomic location (kb)
20,750
21,000
21,250
21,500
21,750
22,00
TTC5
CCNB
PARP2
TEP1
KLHL3
RNASE
ANG
RNASE
RNASE
RNASE
RNASE
HNRNI
RPGRI
SUPT1
CHD8
RAB2E
TOX4
METTL
SALL2
−log10 GWAS p−value