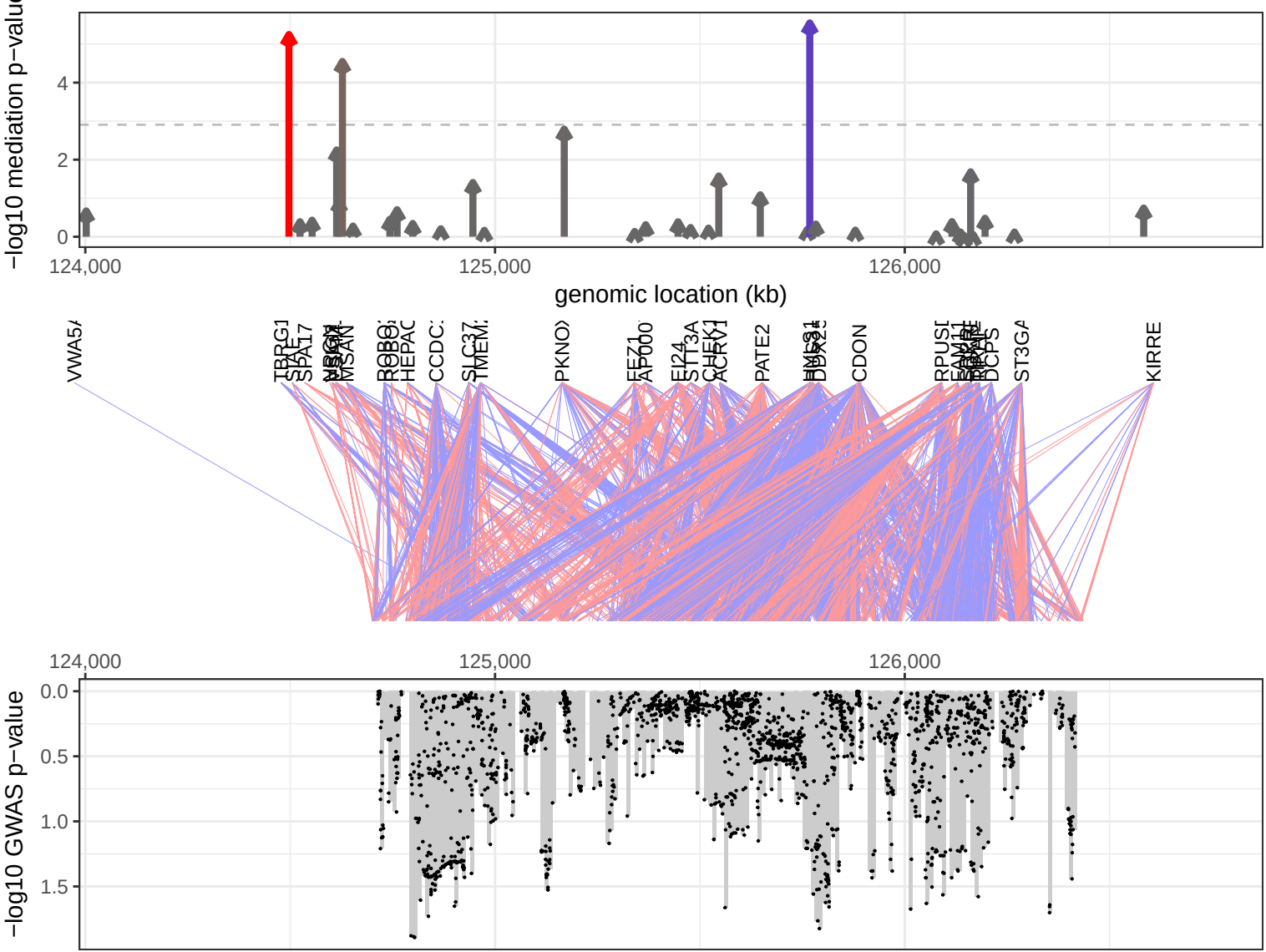
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2
4
124,000
125,000
126,000
genomic location (kb)
VWA5A
TBRG1
SPAT17
MSAN
ROBO
HEPAC
CCDC
SLC37
TMEM
PKNOX
FEZ1
APOO
EI24
ST3A
CHEK1
ACRV1
PATE2
CDON
RPUSD
DCPS
ST3GA
KIRRE
124,000
125,000
126,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5