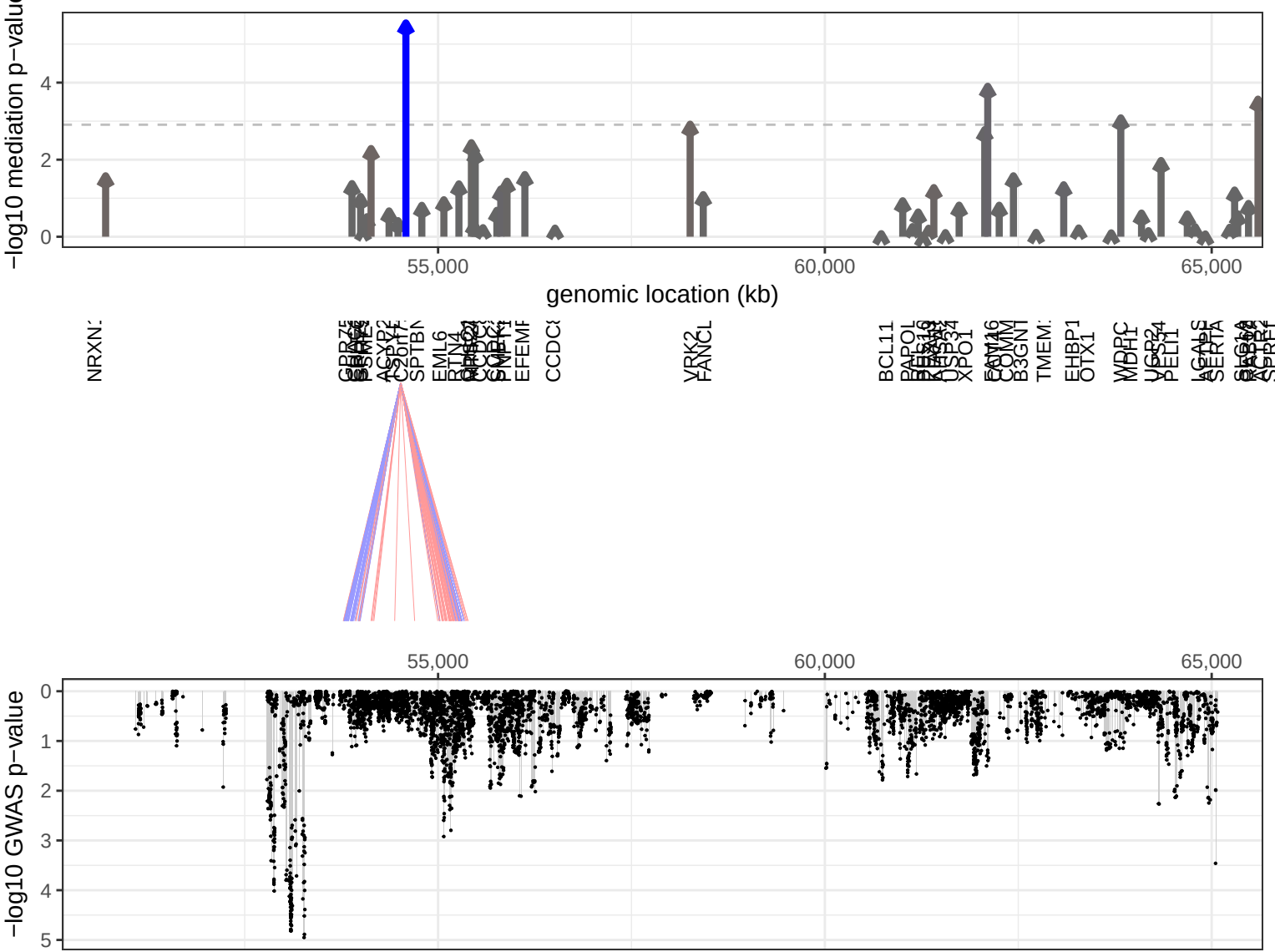
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2
4
55,000
60,000
65,000
genomic location (kb)
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SPTBN
EML6
CCDC
VRK2
FANCL
BCL11
XPO1
B3GNT
TMEM
EHBP1
OTX1
MBHT
PEL1
SERTA
55,000
60,000
65,000
−log10 GWAS p−value
0
1
2
3
4
5