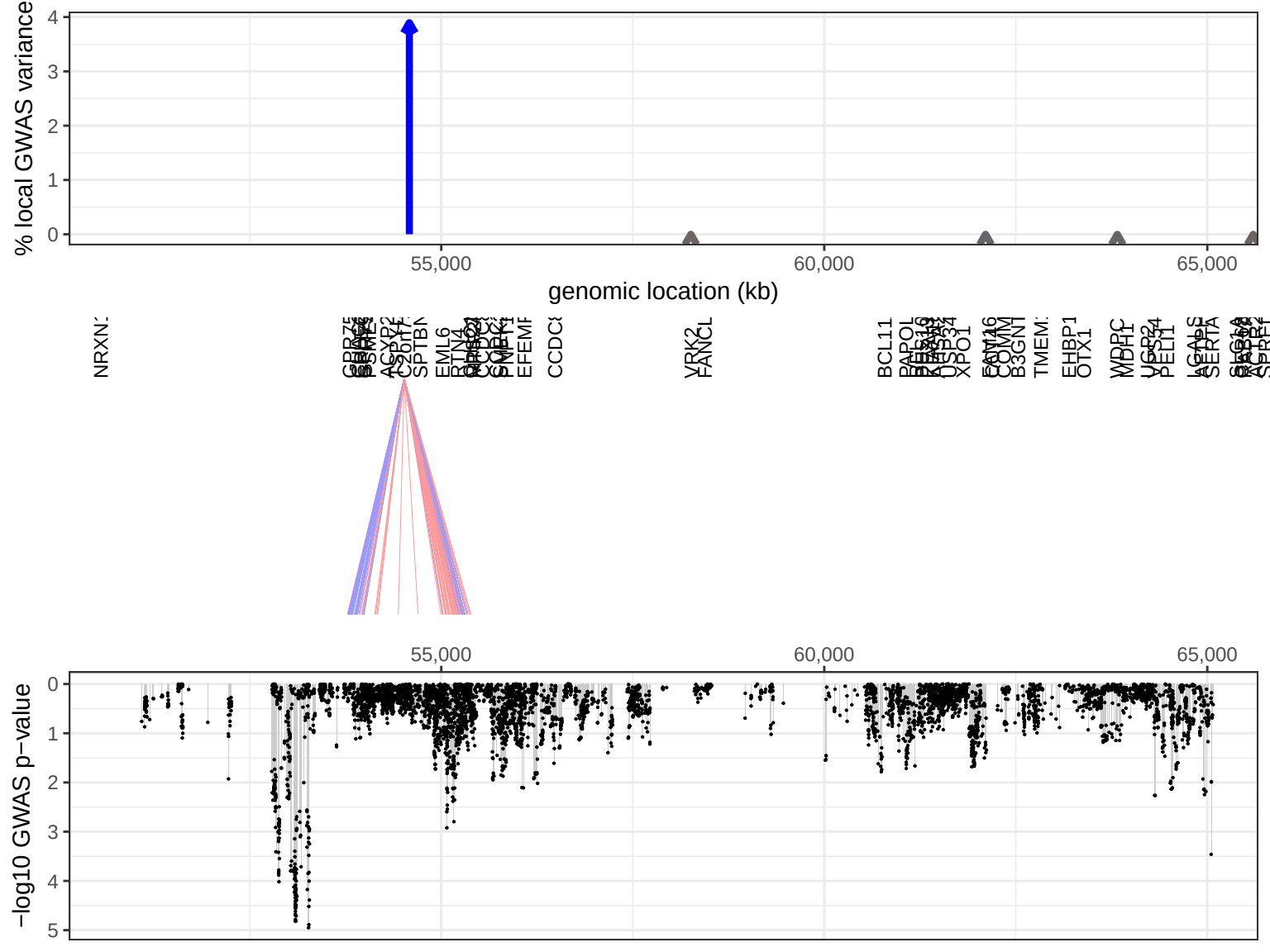
% local GWAS variance
4
3
2
1
0
55,000
60,000
65,000
genomic location (kb)
NRXN
SPTBN
EML6
CCDC
VRK2
FANCL
BCL11
XPO1
B3GNT
TMEM
EHBP1
OTX1
MBHT
PELI1
SERTA
55,000
60,000
65,000
−log10 GWAS p−value
0
1
2
3
4
5