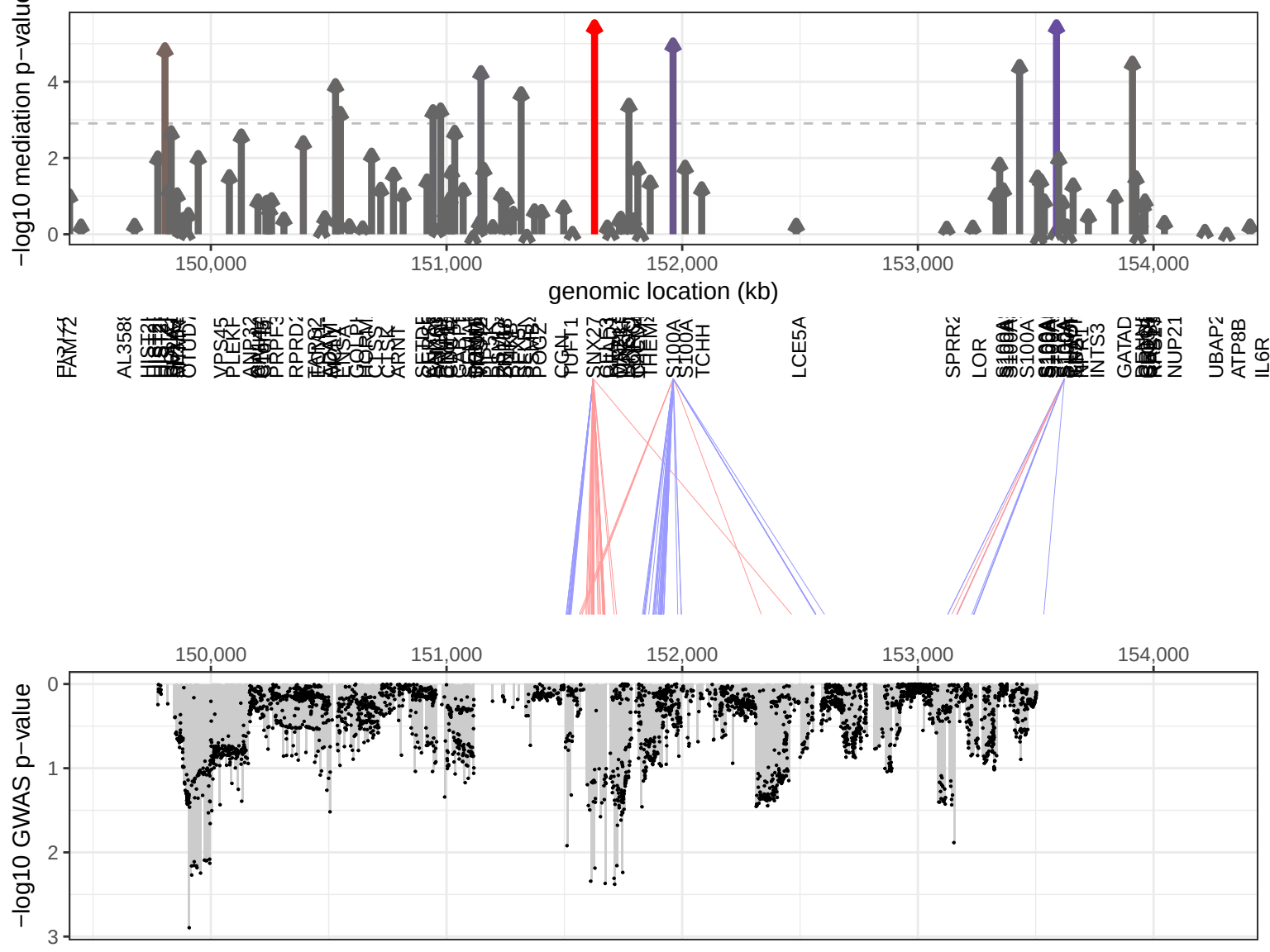
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2
4
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151,000
152,000
153,000
154,000
genomic location (kb)
AL358
VPS45
RPRD2
CGN
TUFT1
SNX27
S100A
TCHH
LCE5A
SPRR2
LOR
S100A
INTS3
GATAD
NUP21
UBAP2
ATP8B
IL6R
−log10 GWAS p−value
0
1
2
3