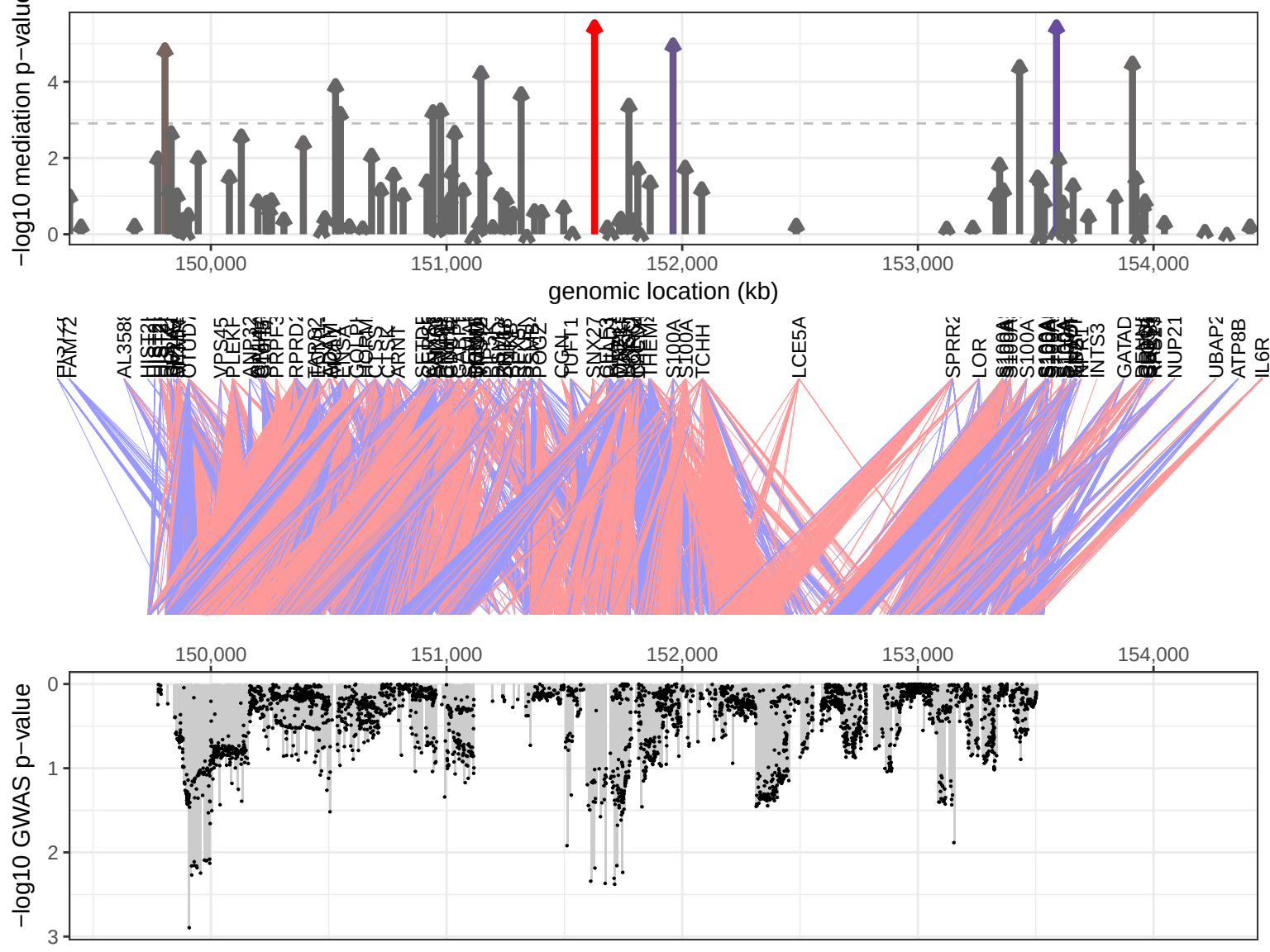
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4
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151,000
152,000
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genomic location (kb)
FAM72
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YPS45
LEKR
RPRD2
CGA
FUT1
SNX27
S100A
TCHH
LCE5A
SPRR2
LOR
S100A
INTS3
GATAD
NUP21
UBAP2
ATP8B
IL6R
150,000
151,000
152,000
153,000
154,000
−log10 GWAS p−value
0
1
2
3