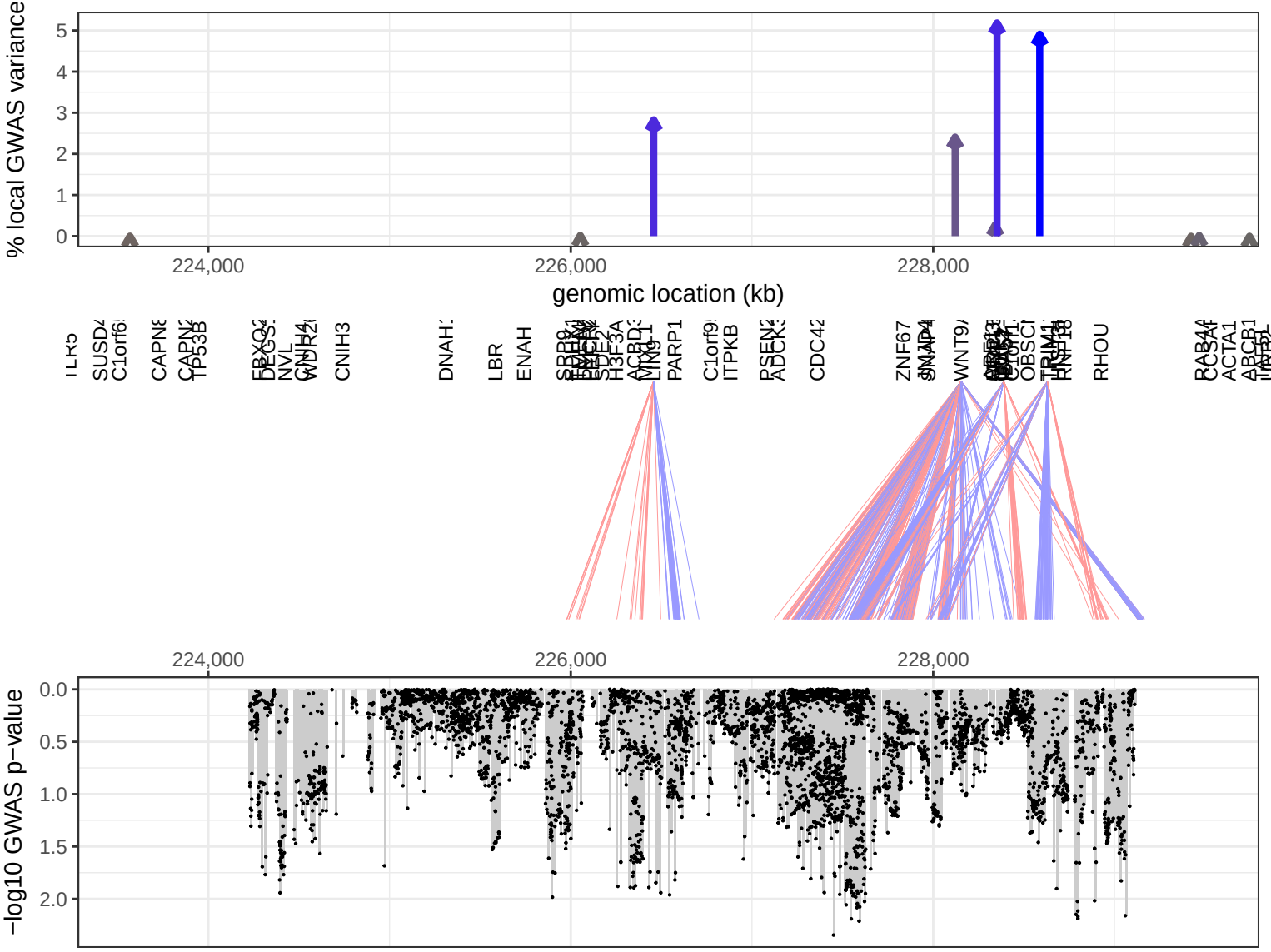
% local GWAS variance
0
1
2
3
4
5
224,000
226,000
228,000
genomic location (kb)
LBR
ENAH
PARP1
ITPKB
CDC42
WNT9A
RHOU
ACTA1
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0
224,000
226,000
228,000