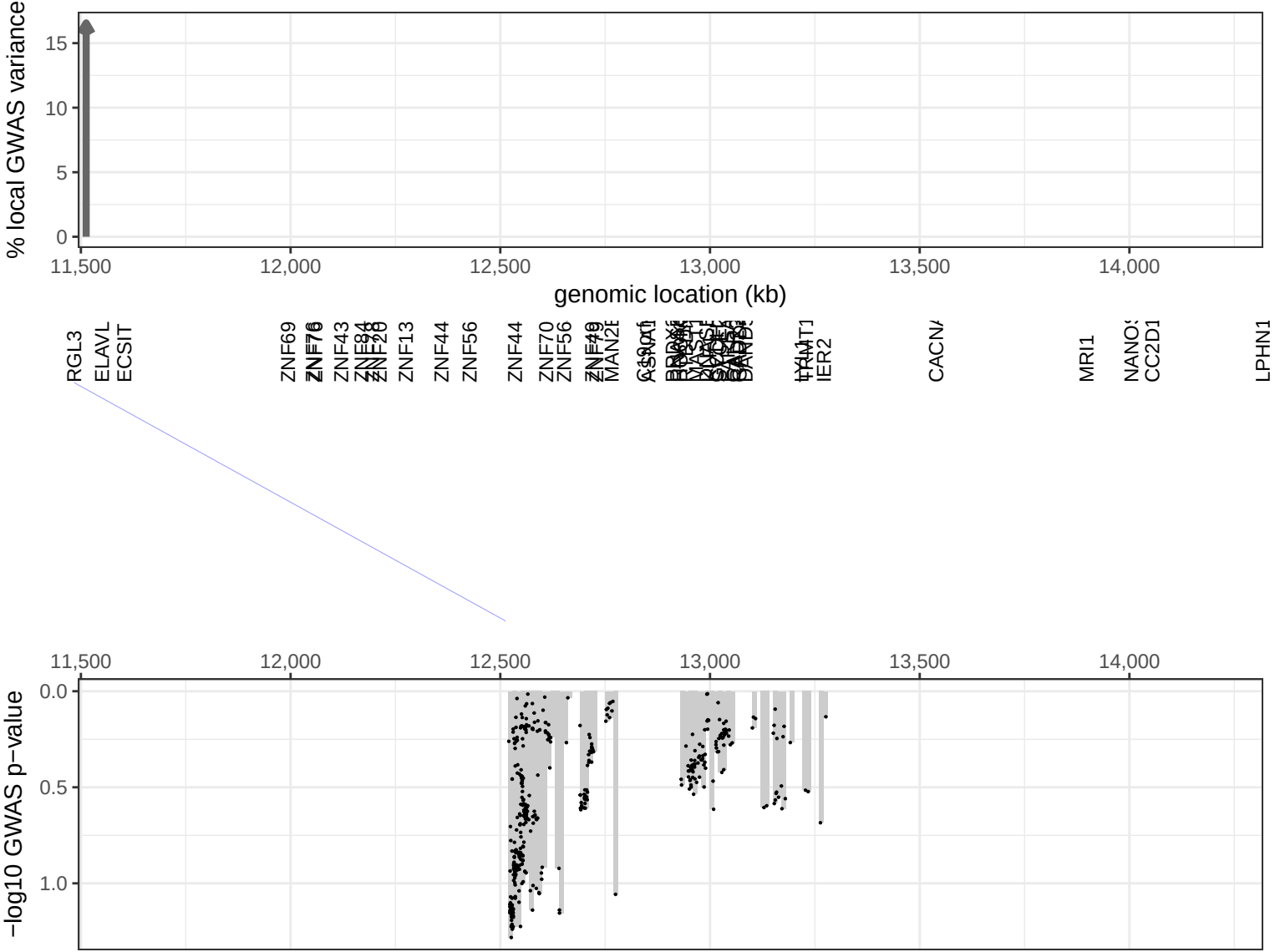
% local GWAS variance
15
10
5
0
11,500
12,000
12,500
13,000
13,500
14,000
genomic location (kb)
RGL3
ELAVL
ECSIT
ZNF69
ZNF43
ZNF13
ZNF44
ZNF56
ZNF44
ZNF70
ZNF56
MAN2E
IER2
MRI1
CC2D1
LPHN1
−log10 GWAS p−value
0.0
0.5
1.0