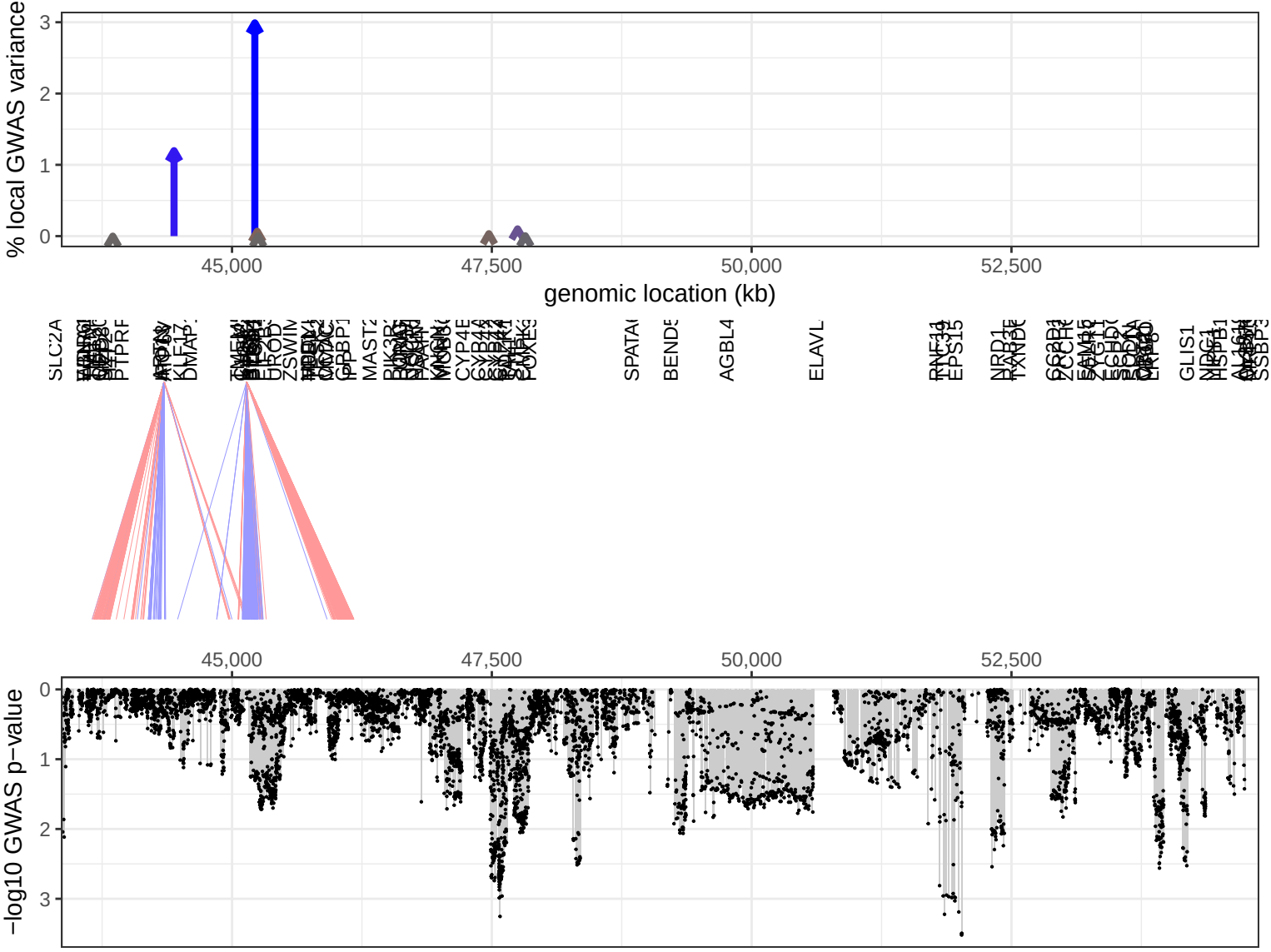
% local GWAS variance
3
2
1
0
45,000
47,500
50,000
52,500
genomic location (kb)
SLC2A
PTPRF
DMAP
MAST2
CYP4E
SPATA
BEND5
AGBL4
ELAVL
EPS15
NRD1
GLIS1
−log10 GWAS p−value
0
1
2
3