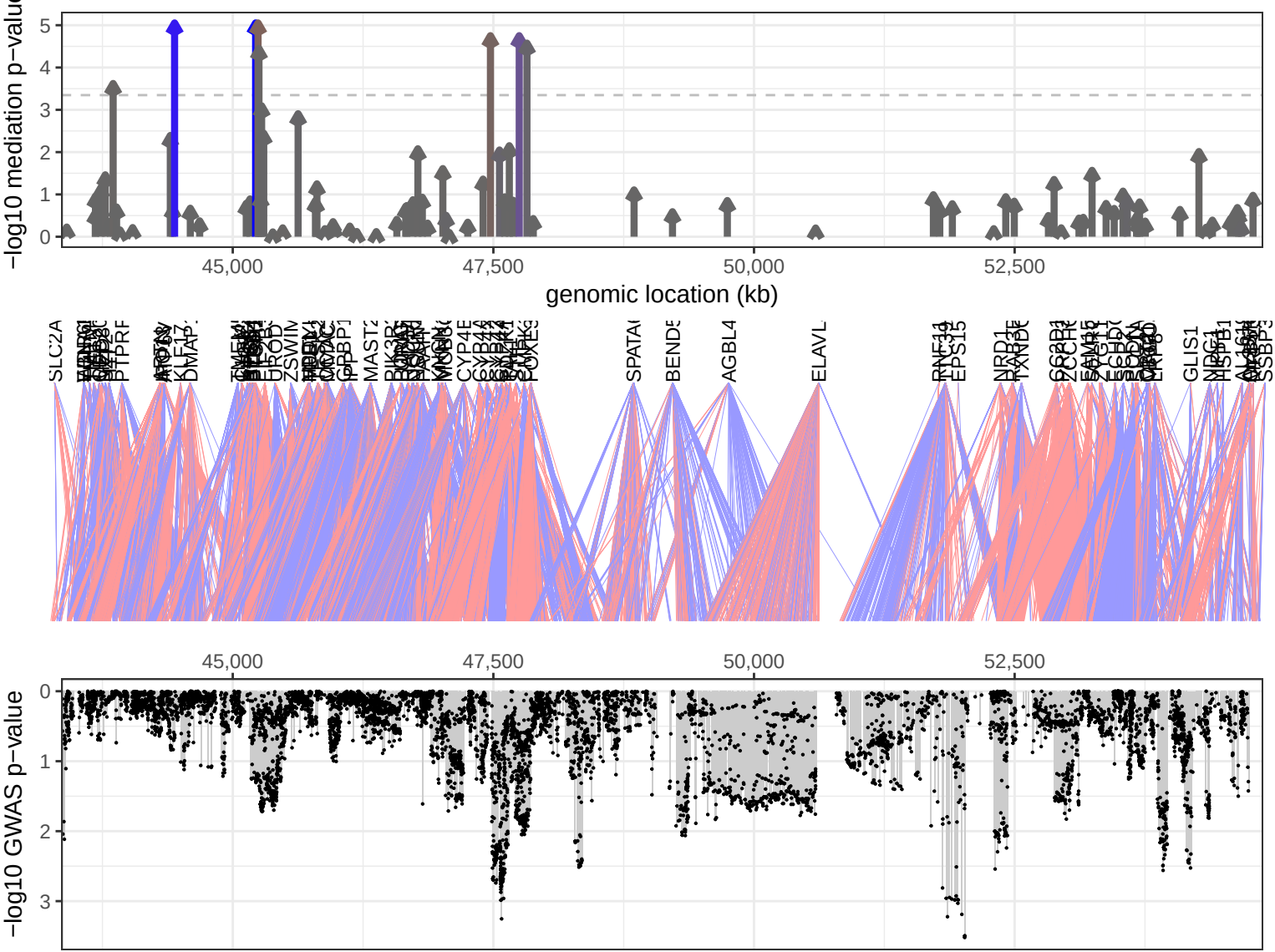
−log10 mediation p−value
genomic location (kb)
45,000
47,500
50,000
52,500
SPATA
BEND
AGBL4
ELAVL
GLIS1
−log10 GWAS p−value