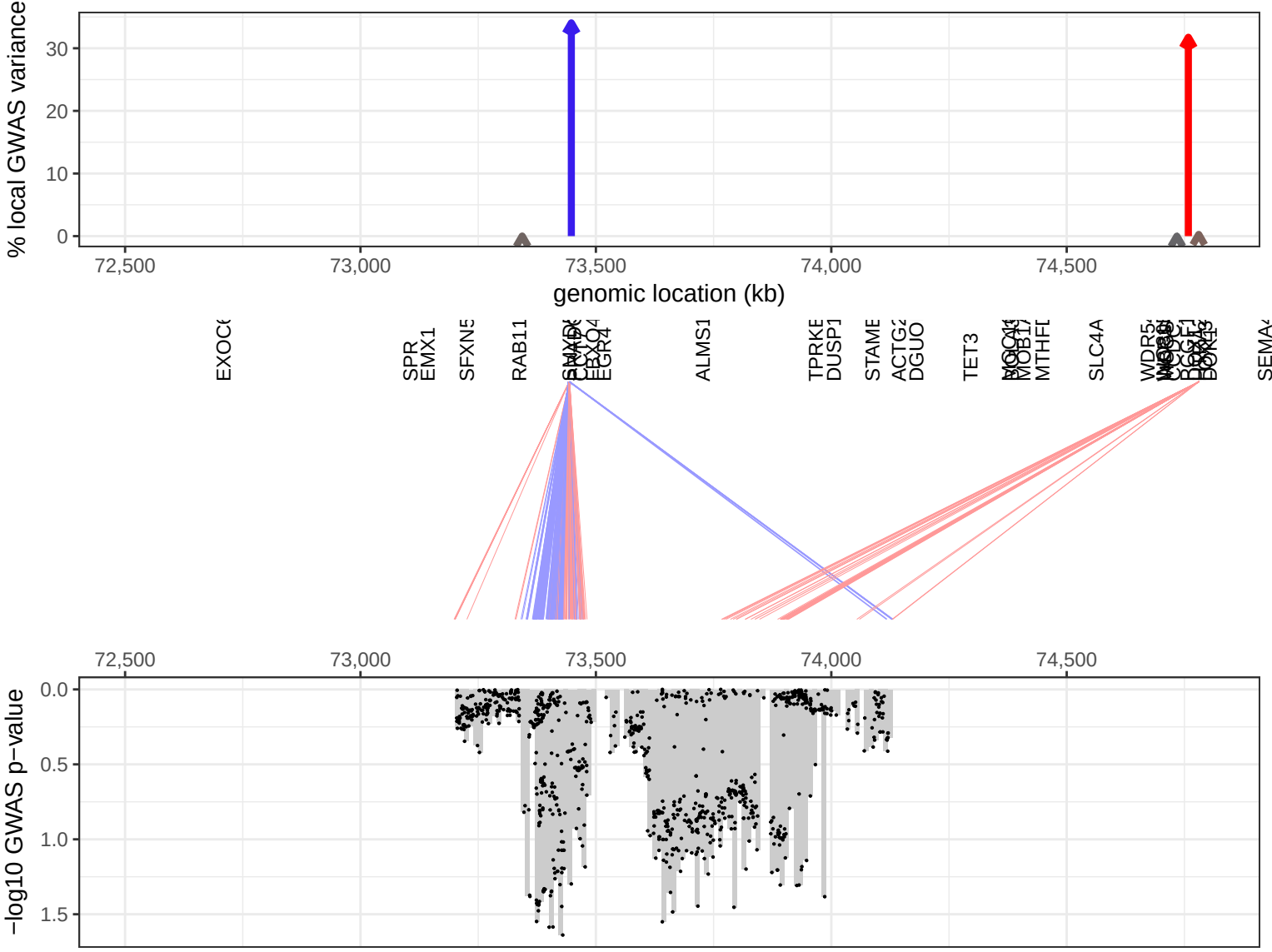
% local GWAS variance
30
20
10
0
72,500
73,000
73,500
74,000
74,500
genomic location (kb)
SPR
EMX1
RAB11
ALMS1
TPRKB
DUSP1
STAME
TET3
SLC4A
−log10 GWAS p−value
0.0
0.5
1.0
1.5