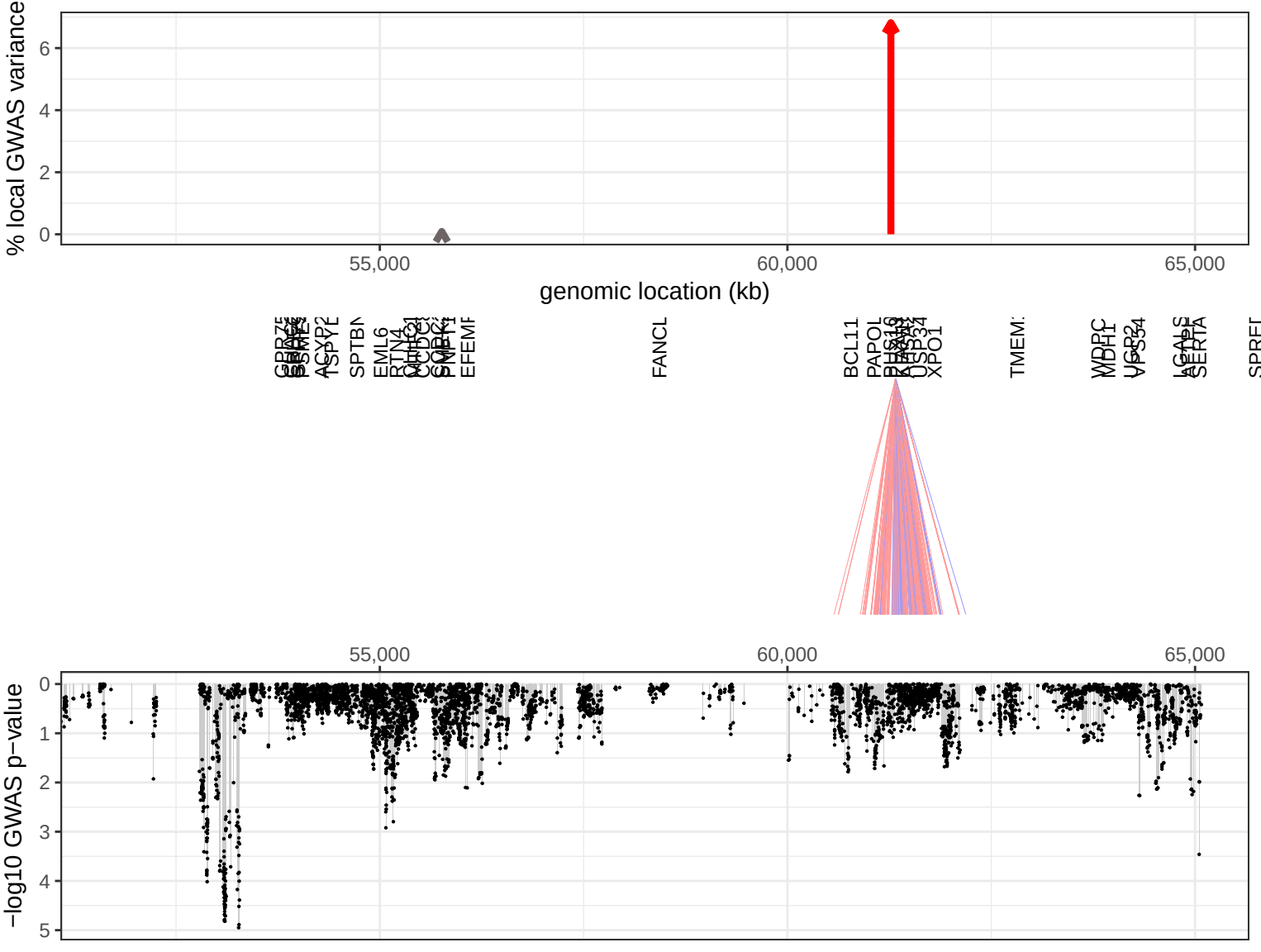
% local GWAS variance
0
2
4
6
55,000
60,000
65,000
genomic location (kb)
SPTBN
EML6
EFEMP
FANCL
BCL11
PAPOL
XPO1
TMEM
SERTA
55,000
60,000
65,000
−log10 GWAS p−value
0
1
2
3
4
5