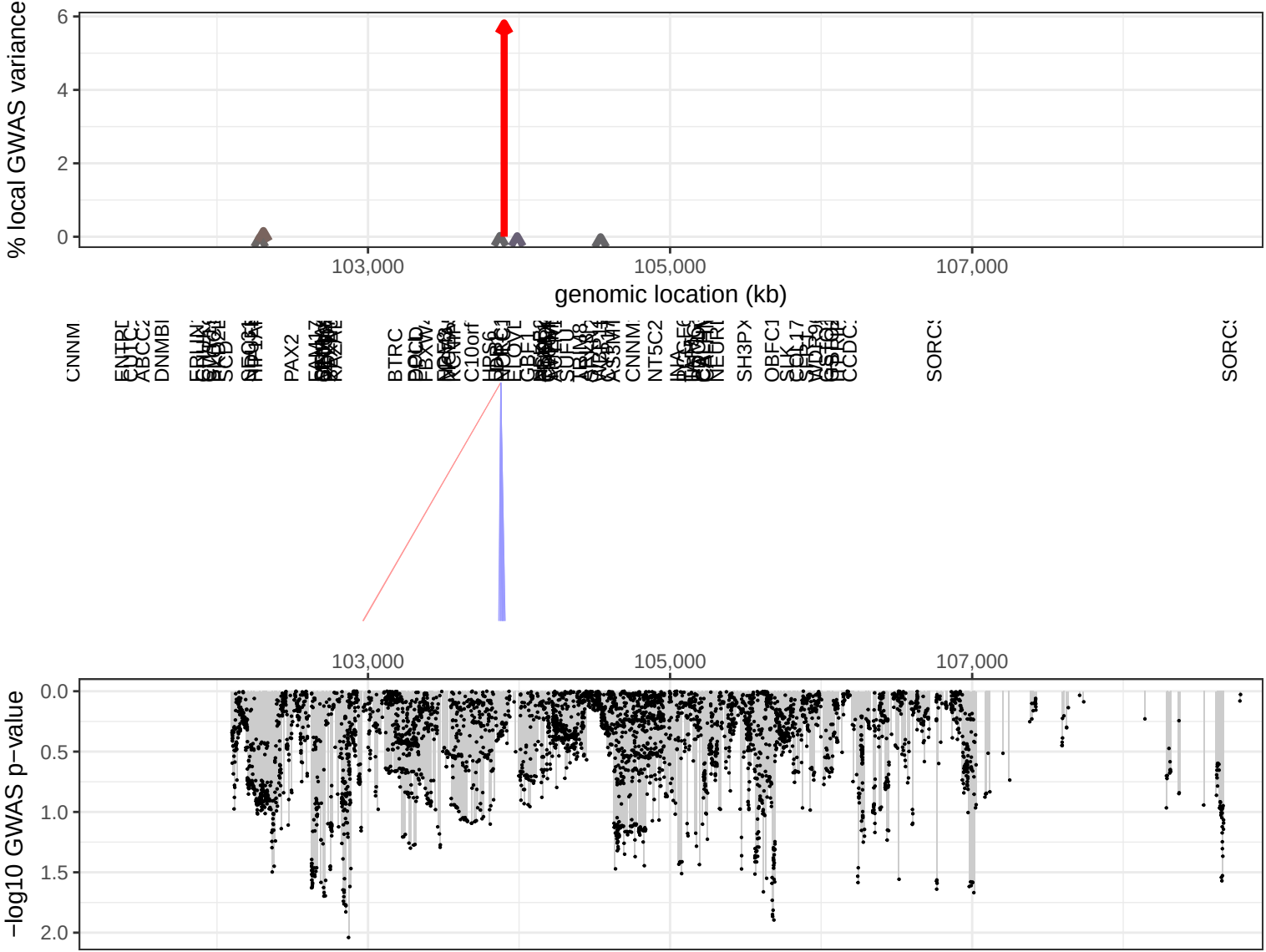
% local GWAS variance
6
4
2
0
103,000
105,000
107,000
genomic location (kb)
CNNM
PAX2
BTRC
C10orf
NT5C2
SH3PX
OBFC1
SORC
SORC
103,000
105,000
107,000
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0