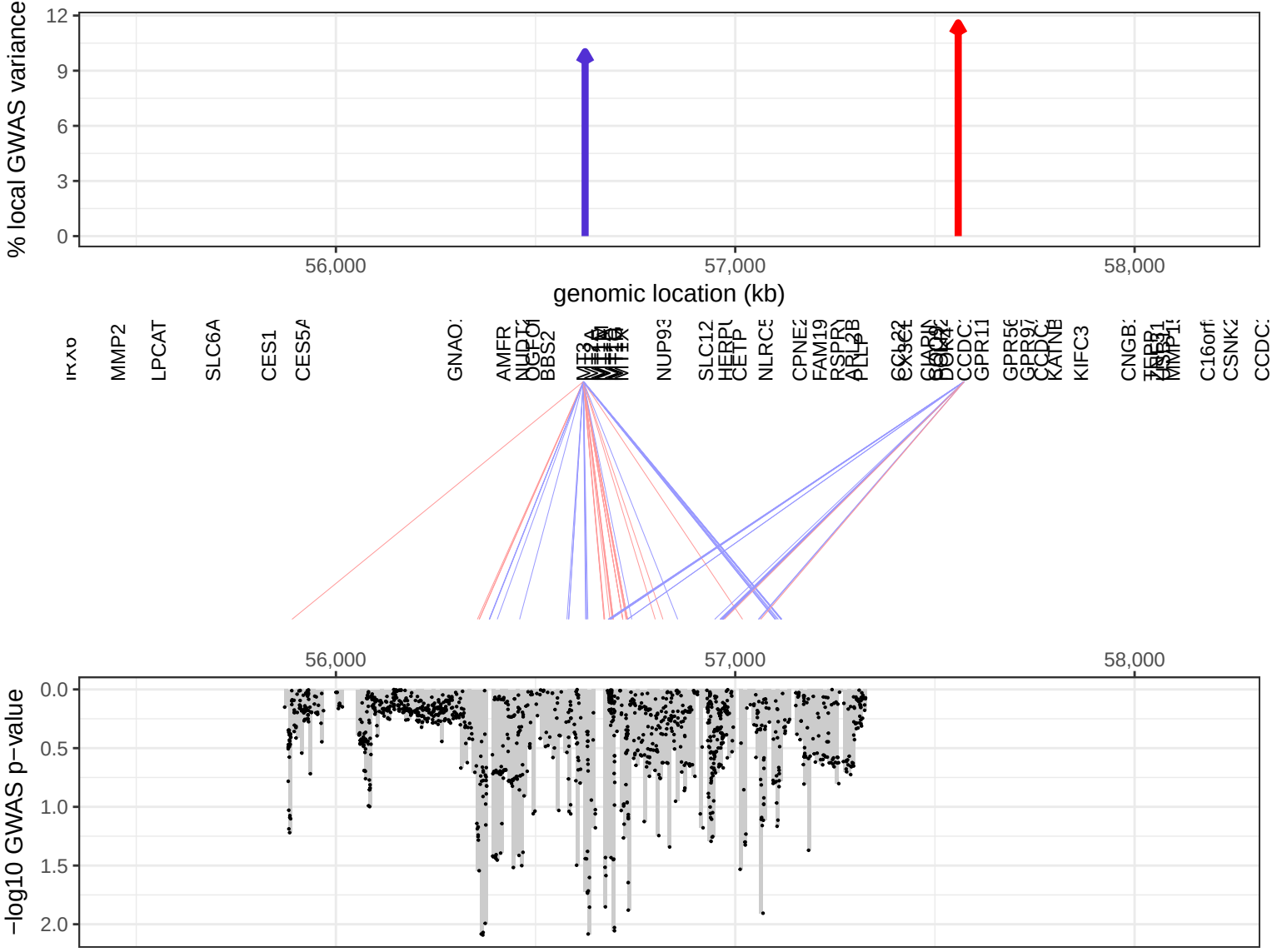
% local GWAS variance
12
9
6
3
0
56,000
57,000
58,000
genomic location (kb)
MMP2
LPCAT
SLC6A
CES1
BBS2
NUP93
SLC12
CETP
NLRC5
CPNE2
FAM19
GPR11
KIFC3
CNGB1
C16orf
CSNK2
−log10 GWAS p−value
0.0
0.5
1.0
1.5
2.0