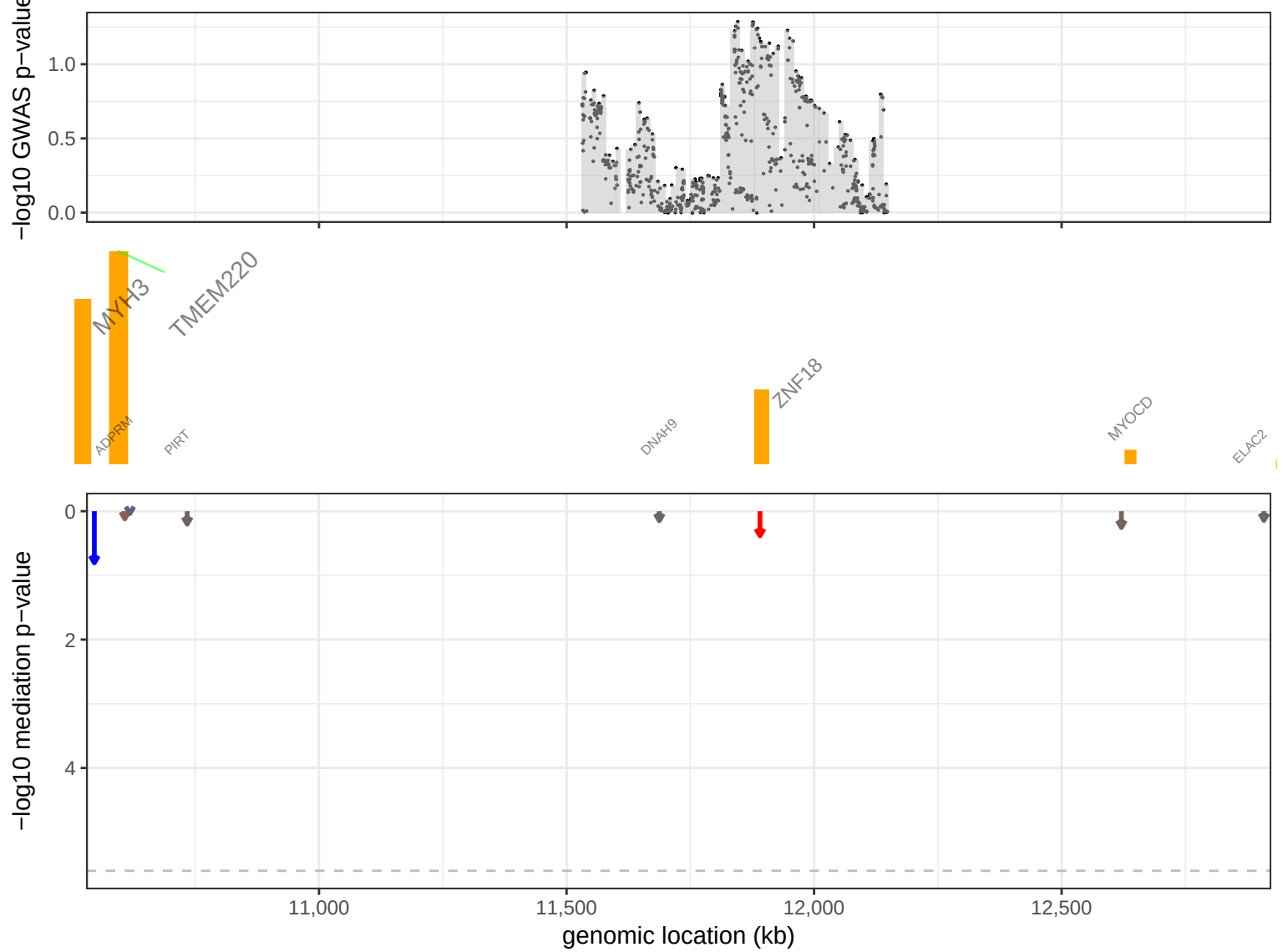
−log10 GWAS p−value
1.0
0.5
0.0
MYH3
TMEM220
ADPRM
PIRT
DNAH9
ZNF18
MYOCD
ELAC2
−log10 mediation p−value
0
2
4
11,000
11,500
12,000
12,500
genomic location (kb)