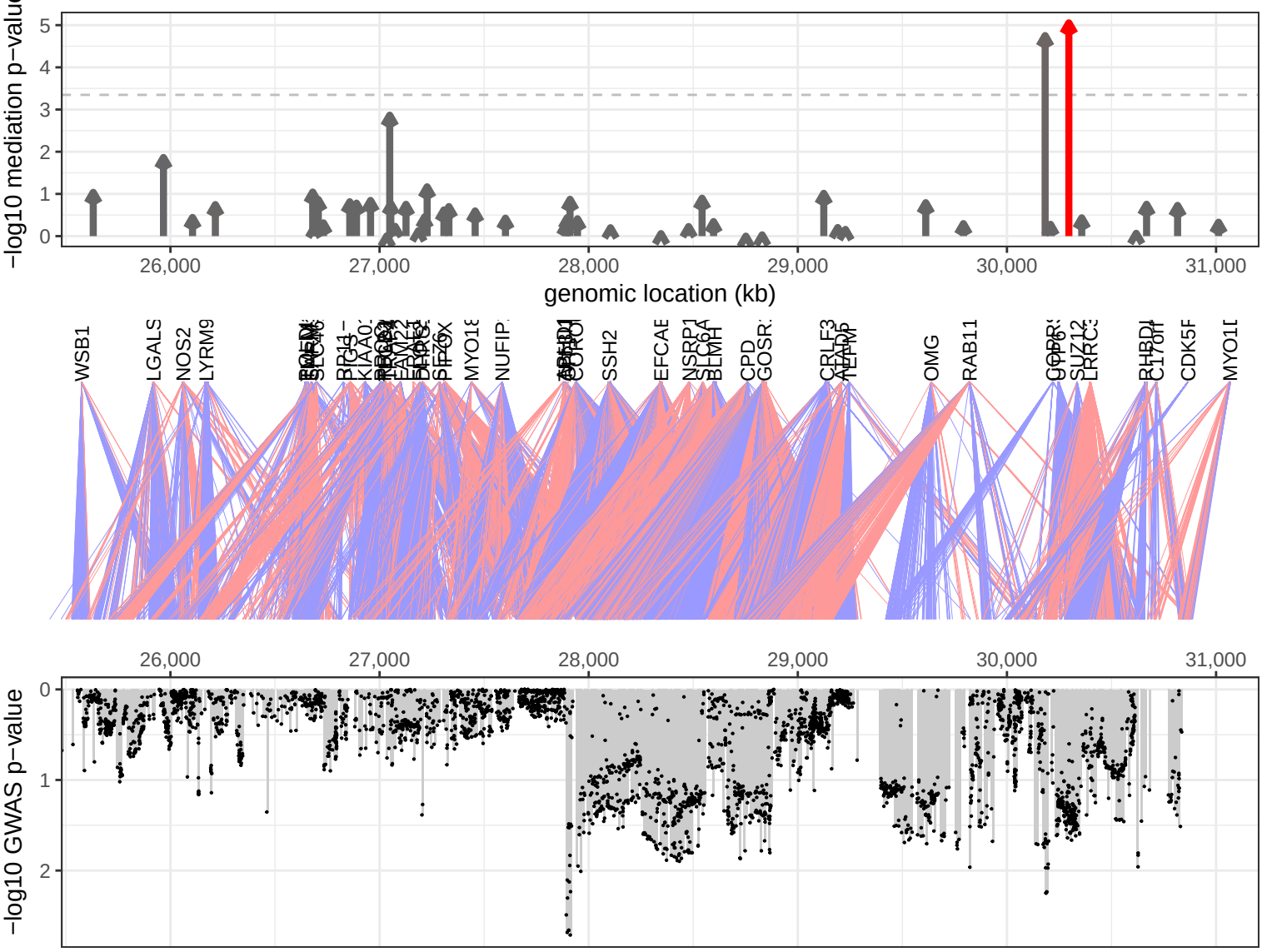
−log10 mediation p−value
genomic location (kb)
26,000
27,000
28,000
29,000
30,000
31,000
WSB1
LGALS
NOS2
LYRM9
MYO1E
NUFIP
SSH2
EFCAE
CPD
OMG
RAB11
CDK5F
−log10 GWAS p−value