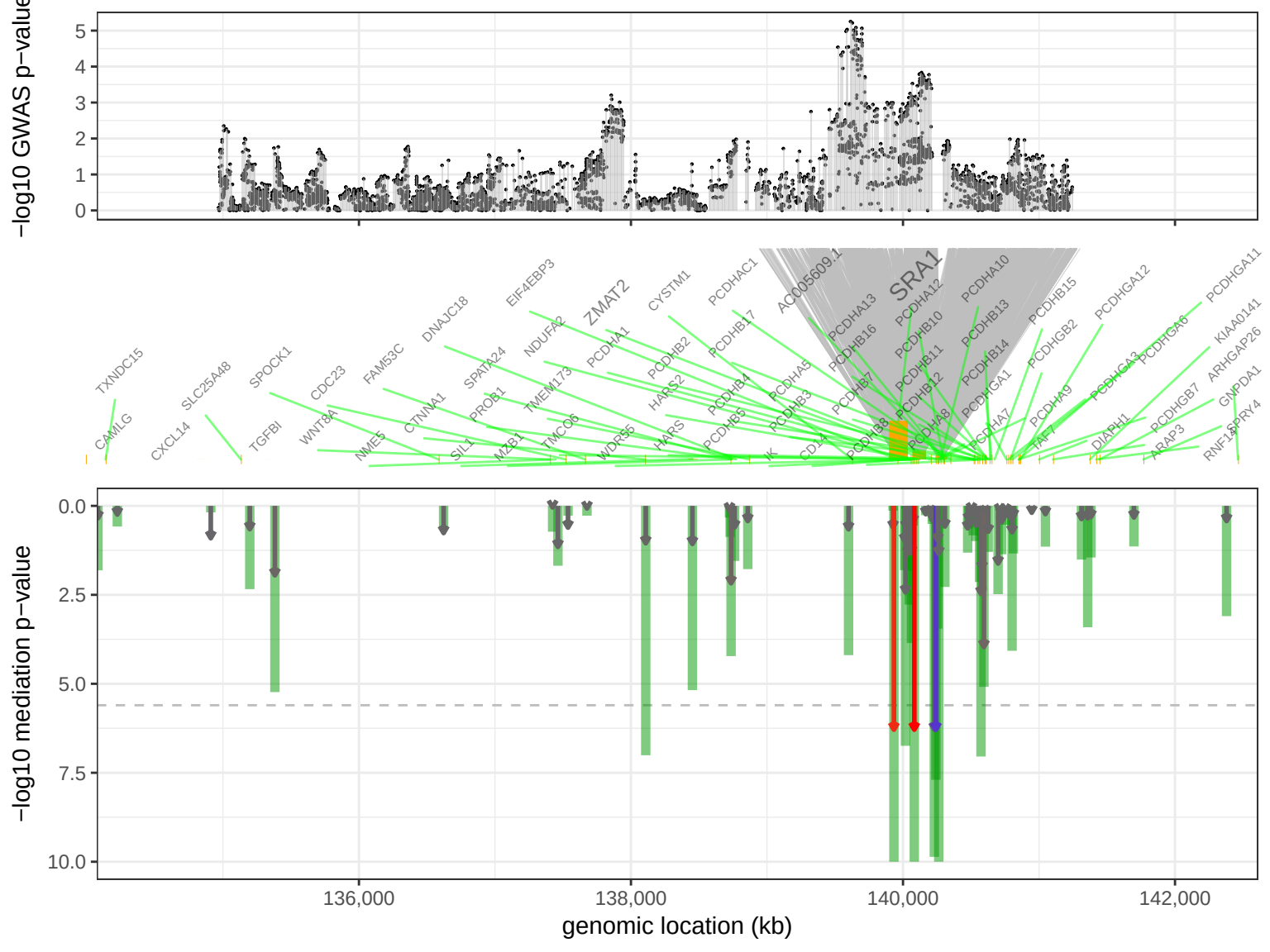
−log10 GWAS p−value
5
4
3
2
1
0
TXNDC15
CAMLG
CXCL14
SLC25A48
TGFBI
SPOCK1
WNT8A
CDC23
NME5
FAM53C
CTNNA1
SIL1
DNAJC18
MZB1
SPATA24
PROB1
EIF4EBP3
TMCO6
TMEM173
NDUFA2
WDR95
ZMAT2
HARS
PCDHA1
HARS2
CYSTM1
PCDHB2
PCDHB5
PCDHAC1
PCDHB4
PCDHB17
PCDHA5
PCDHB3
AC005609.1
PCDHB7
IK
CDA
PCDHA13
PCDHB16
PCDHB8
SRA1
PCDHA12
PCDHB10
PCDHB11
PCDHB12
PCDHB48
PCDHA10
PCDHB13
PCDHB14
PCDHGA1
PCDHA7
PCDHB15
PCDHGB2
PCDHA9
TAF7
PCDHGA12
PCDHGA3
PCDHGA6
DIAPH1
PCDHGB7
ARAP3
PCDHGA11
KIAA0141
ARHGAP26
GMPDA1
RNF14
SPRY4
−log10 mediation p−value
0.0
2.5
5.0
7.5
10.0
136,000
138,000
140,000
142,000
genomic location (kb)