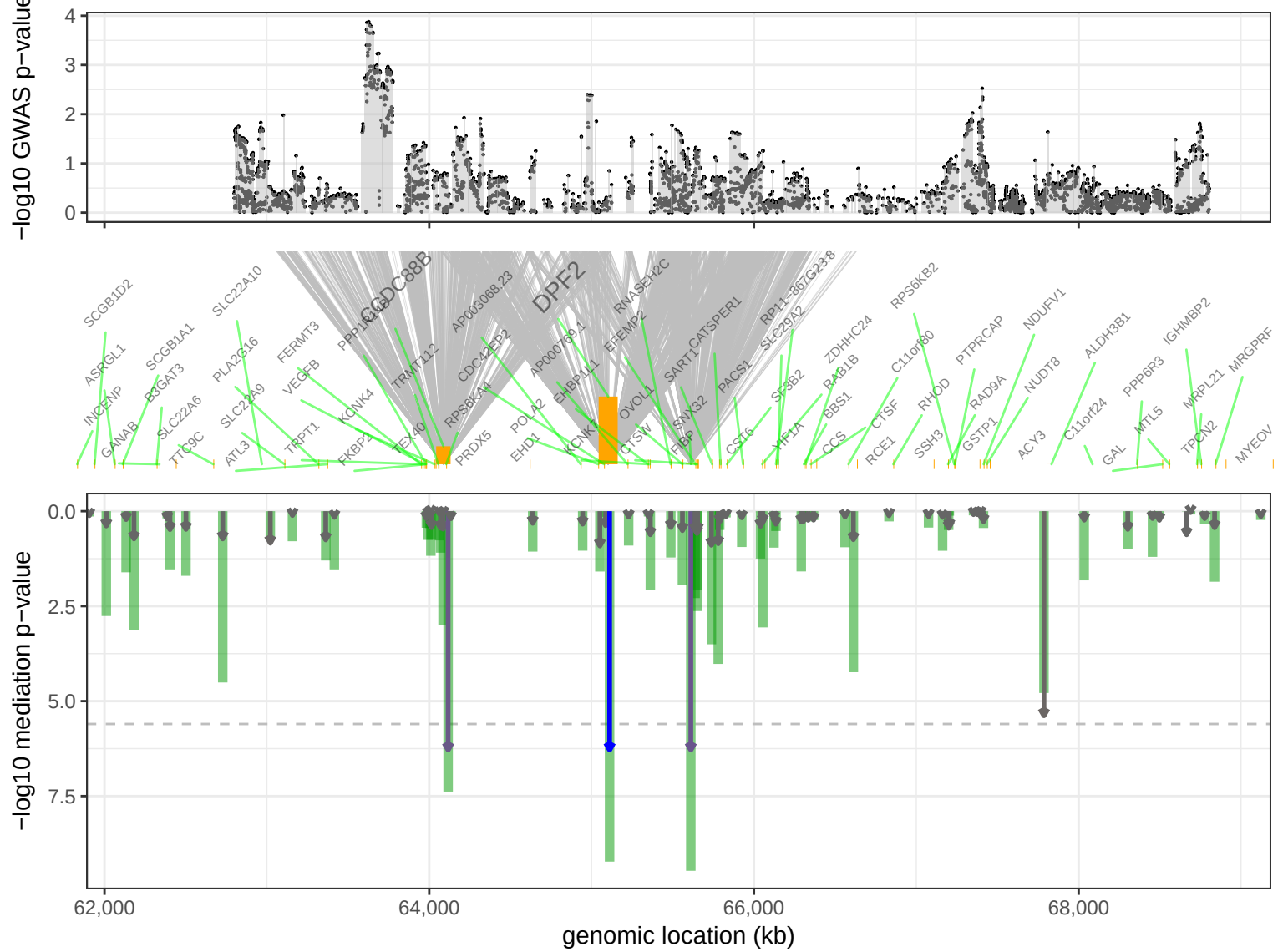
−log10 GWAS p−value
4
3
2
1
0
GGDC88B
DPF2
SCGB1D2
ASRGL1
INCENP
GANAB
SCGB1A1
B3GAT3
SLC22A6
TTC9C
SLC22A10
PLA2G16
SLC22A9
ATL3
FERMT3
VEGFB
TRPT1
KCNK4
FKBP2
PPP1R14B
TRMT112
TEX40
AP003068.23
CDC42EP2
RPS6KA1
PRDX5
POLR2
EHD1
AP000769.1
EHBP1L1
KCNK7
RNASEH2C
EFEMP2
OVOL1
CTSW
SART1
CATSPER1
SNX32
FIBP
PACS1
CSF6
RP11-867G23.8
SLC29A2
SF3B2
HIF1A
ZDHHC24
RAB1B
BBS1
CCS
RPS6KB2
C11orf80
CTSF
RCE1
RHOD
SSH3
PTPRCAP
RAD9A
GSTP1
NDUFV1
NUDT8
ACY3
ALDH3B1
C11orf24
GAL
PPP6R3
MTL5
IGHMBP2
MRPL21
TPCN2
MRGPRF
MYEOV
−log10 mediation p−value
0.0
2.5
5.0
7.5
62,000
64,000
66,000
68,000
genomic location (kb)