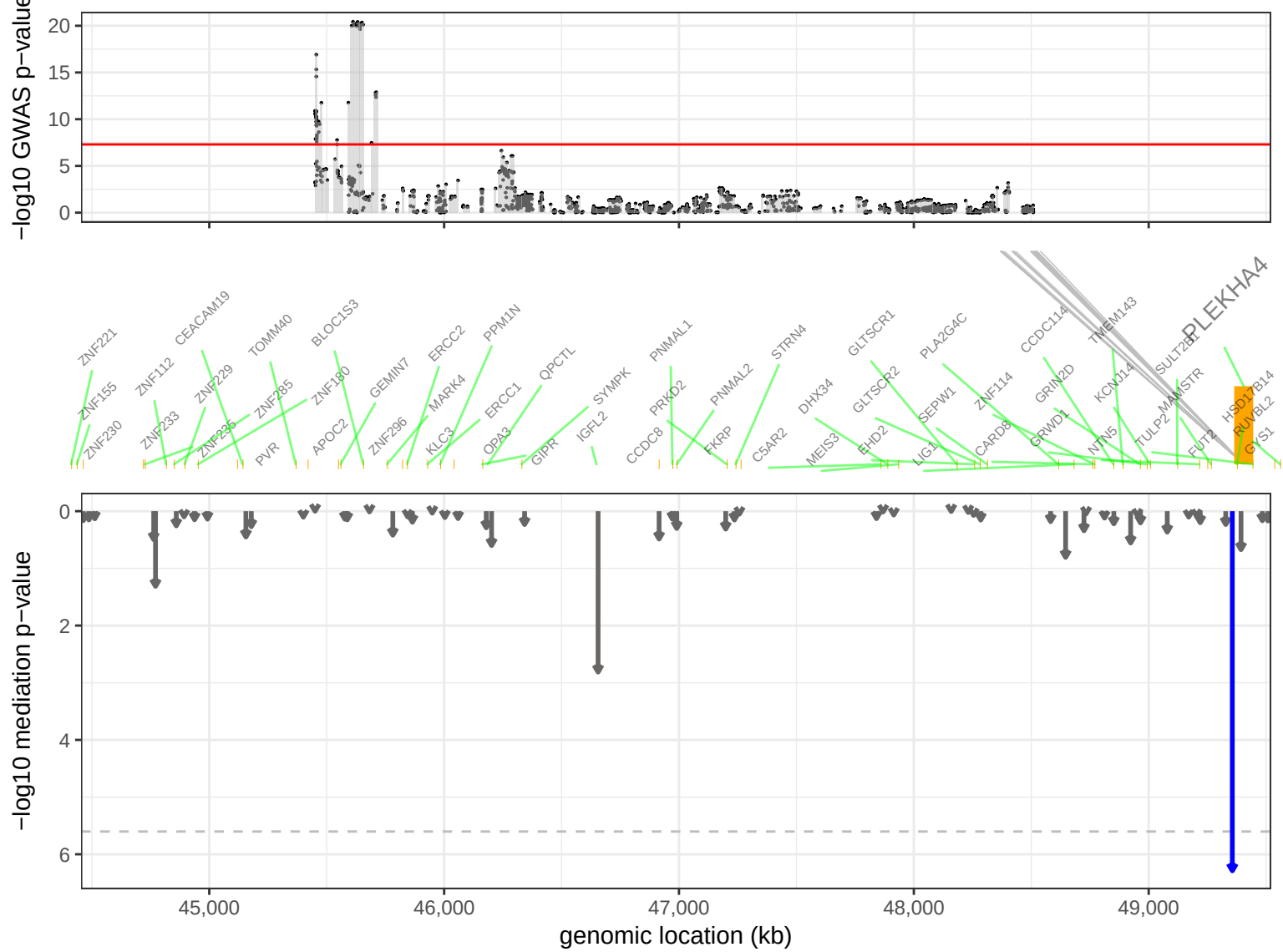
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15
10
5
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ZNF230
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TOMM40
ZNF285
PVR
BLOC1S3
ZNF100
APOC2
GEMIN7
ZNF296
ERCC2
MARK4
KLC3
PPM1N
ERCC1
OPA3
QPCTL
GIPR
SYMPK
IGFL2
PNMAL1
PRKD2
CCDC8
PNMAL2
FKRP
STRN4
C5AR2
DHX34
MEIS3
GLTSCR1
GLTSCR2
EHD2
PLA2G4C
SEPW1
LIG4
ZNF114
CARD8
CCDC114
GRIN2D
GRWD1
TMEM143
KCNJ14
NTN5
SULT2B1
TULP2
MAMSTR
FUT2
PLEKHA4
HSD17B14
RUVBL2
GYS1
−log10 mediation p−value
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4
6
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46,000
47,000
48,000
49,000
genomic location (kb)