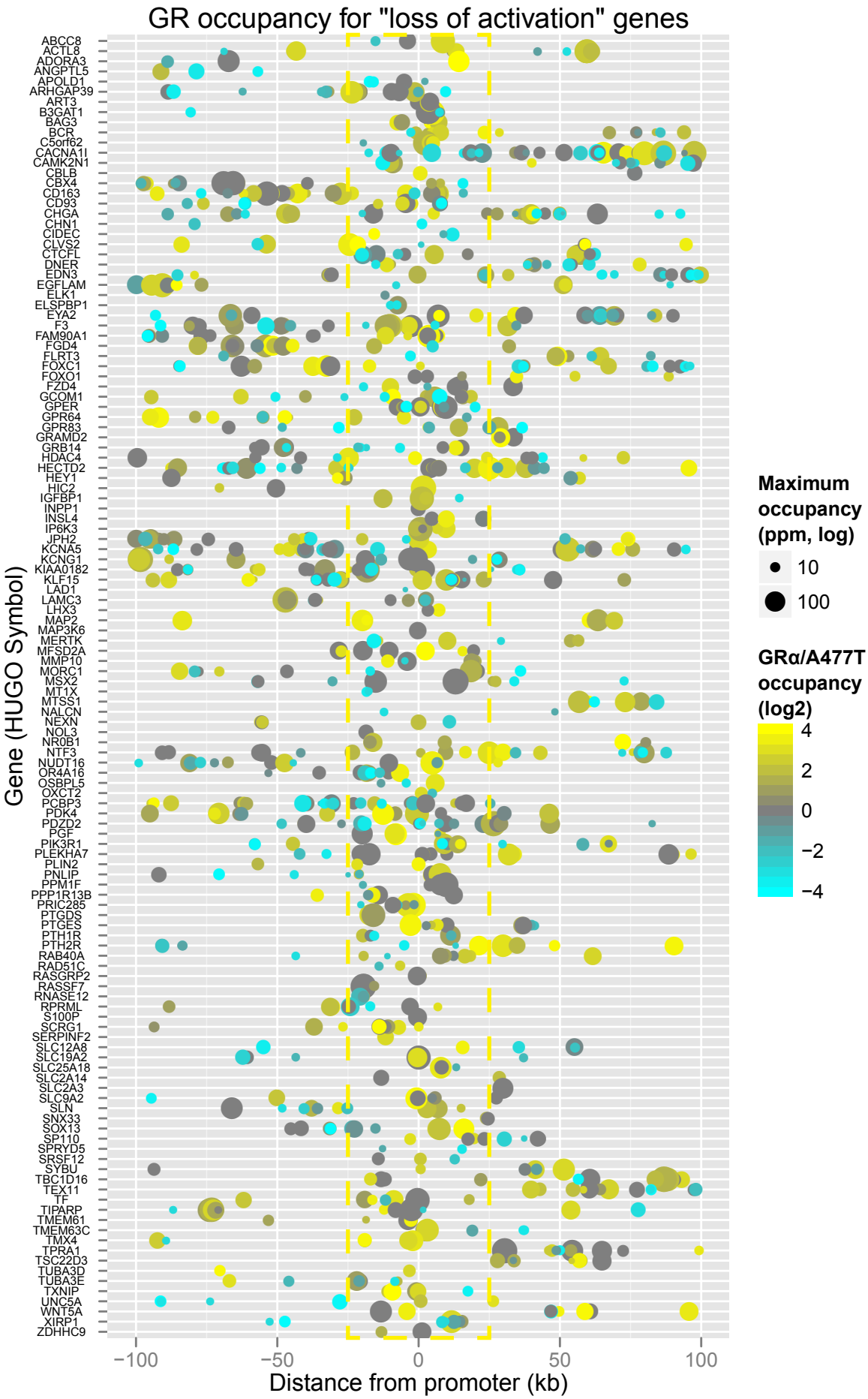
GR occupancy for "loss of activation" genes
Gene (HUGO Symbol)
ABCC8
ACTL8
ADORA3
ANGPTL5
APOLD1
ARHGAP39
ART3
B3GAT1
BAG3
BCR
C5orf62
CACNA1I
CAMK2N1
CBLB
CBX4
CD163
CD93
CHGA
CHN1
CIDEC
CLVS2
CTCFL
DNER
EDN3
EGFLAM
ELK1
ELSPBP1
EYA2
F3
FAM90A1
FGD4
FLRT3
FOXC1
FOXO1
FZD4
GCOM1
GPER
GPR64
GPR83
GRAMD2
GRB14
HDAC4
HECTD2
HEY1
HIC2
IGFBP1
INPP1
INSL4
IP6K3
JPH2
KCNA5
KCNG1
KIAA0182
KLF15
LAD1
LAMC3
LHX3
MAP2
MAP3K6
MERTK
MFSD2A
MMP10
MORC1
MSX2
MT1X
MTSS1
NALCN
NEXN
NOL3
NR0B1
NTF3
NUDT16
OR4A16
OSBPL5
OXCT2
PCBP3
PDK4
PDZD2
PGF
PIK3R1
PLEKHA7
PLIN2
PNLIP
PPM1F
PPP1R13B
PRIC285
PTGDS
PTGES
PTH1R
PTH2R
RAB40A
RAD51C
RASGRP2
RASSF7
RNASE12
RPRML
S100P
SCRG1
SERPINF2
SLC12A8
SLC19A2
SLC25A18
SLC2A14
SLC2A3
SLC9A2
SLN
SNX33
SOX13
SP110
SPRYD5
SRSF12
SYBU
TBC1D16
TEX11
TF
TIPARP
TMEM61
TMEM63C
TMX4
TPRA1
TSC22D3
TUBA3D
TUBA3E
TXNIP
UNC5A
WNT5A
XIRP1
ZDHHC9
−100
−50
0
50
100
Distance from promoter (kb)
Maximum occupancy (ppm, log)
10
100
GRα/A477T occupancy (log2)
4
2
0
−2
−4