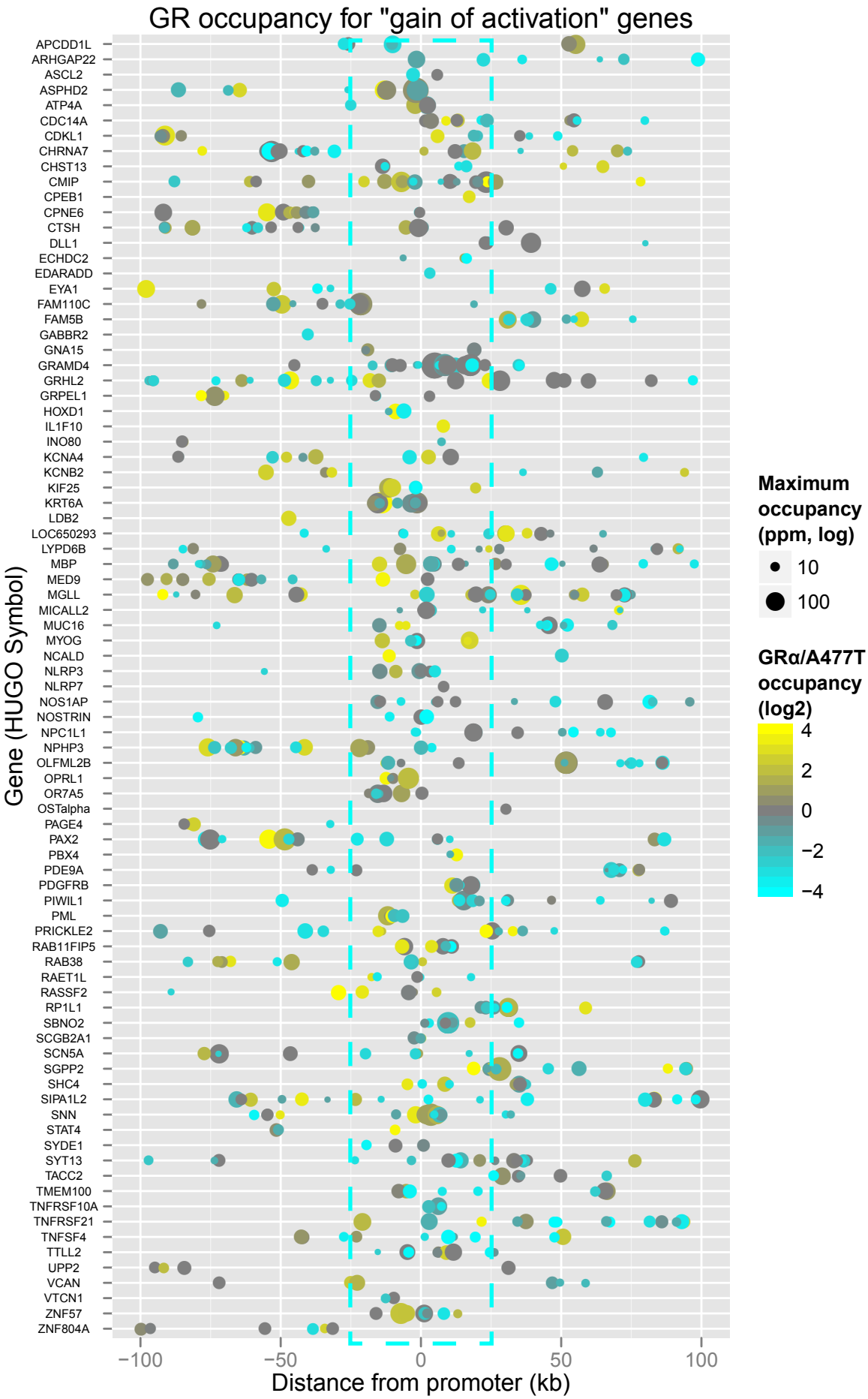
GR occupancy for "gain of activation" genes
Gene (HUGO Symbol)
APCDD1L
ARHGAP22
ASCL2
ASPHD2
ATP4A
CDC14A
CDKL1
CHRNA7
CHST13
CMIP
CPEB1
CPNE6
CTSH
DLL1
ECHDC2
EDARADD
EYA1
FAM110C
FAM5B
GABBR2
GNA15
GRAMD4
GRHL2
GRPEL1
HOXD1
IL1F10
INO80
KCNA4
KCNB2
KIF25
KRT6A
LDB2
LOC650293
LYPD6B
MBP
MED9
MGLL
MICALL2
MUC16
MYOG
NCALD
NLRP3
NLRP7
NOS1AP
NOSTRIN
NPC1L1
NPHP3
OLFML2B
OPRL1
OR7A5
OSTalpha
PAGE4
PAX2
PBX4
PDE9A
PDGFRB
PIWIL1
PML
PRICKLE2
RAB11FIP5
RAB38
RAET1L
RASSF2
RP1L1
SBNO2
SCGB2A1
SCN5A
SGPP2
SHC4
SIPA1L2
SNN
STAT4
SYDE1
SYT13
TACC2
TMEM100
TNFRSF10A
TNFRSF21
TNFSF4
TTLL2
UPP2
VCAN
VTCN1
ZNF57
ZNF804A
−100
−50
0
50
100
Distance from promoter (kb)
Maximum occupancy (ppm, log)
10
100
GRα/A477T occupancy (log2)
4
2
0
−2
−4