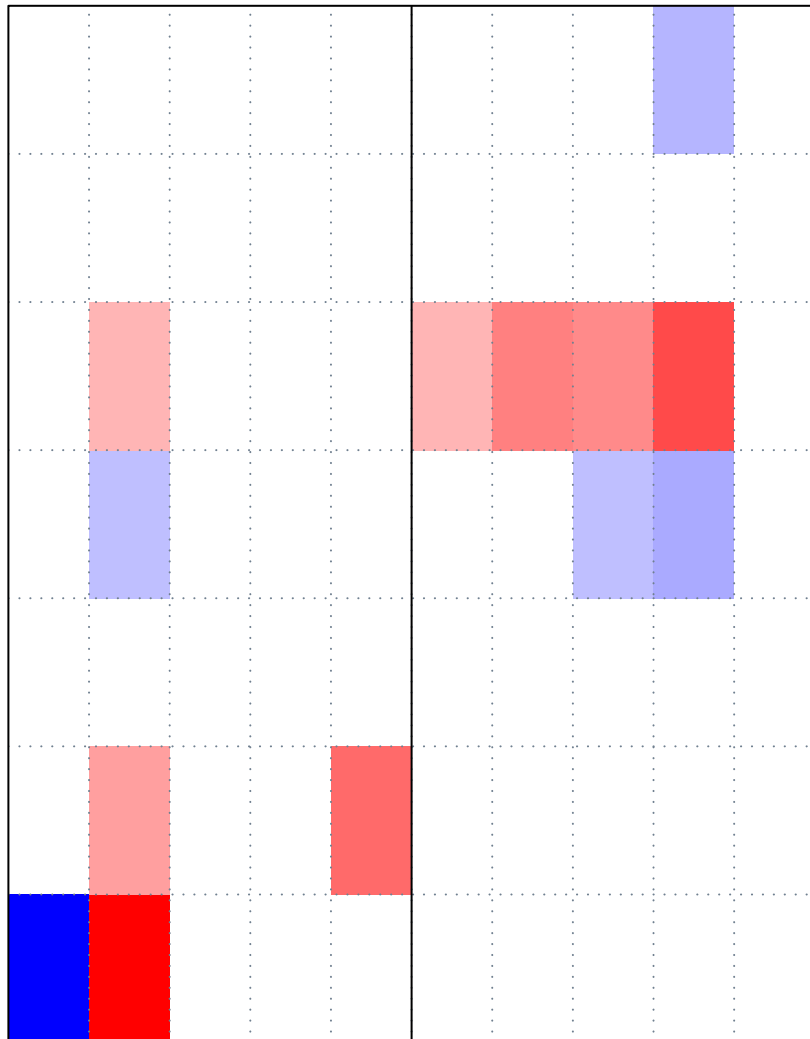


NeuroB

NeuroB_MYC+



ras UP – pmid: 16273092 NA

ras DN – pmid: 16273092 NA

myc UP – pmid: 16273092 NA

myc DN – pmid: 16273092 NA

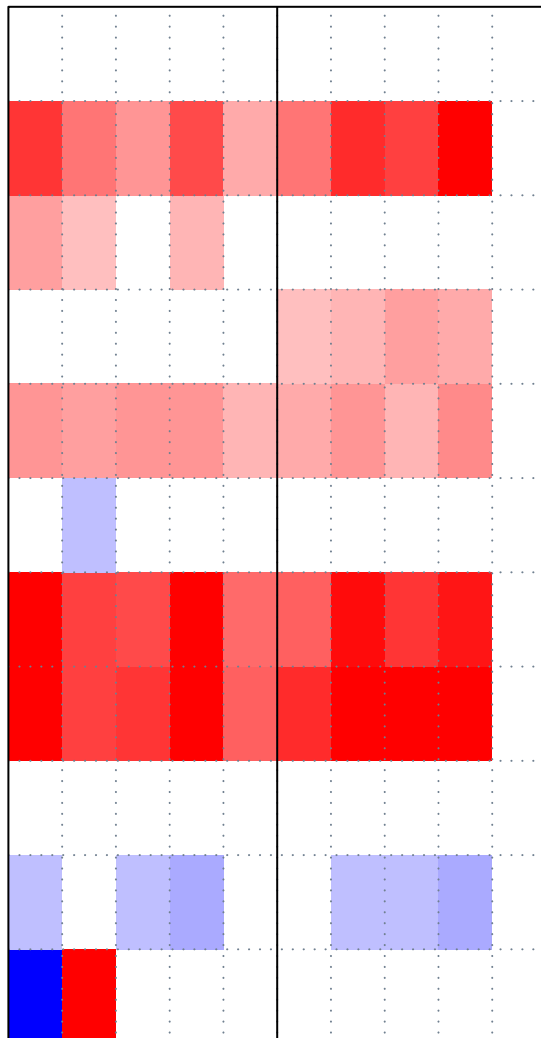
5p NA

5q NA

GS

NeuroB

NeuroB_MYC+



GO:0000041 transition metal ion transport

GO:0000075 cell cycle checkpoint

GO:0000079 regulation of cyclin-dependent protein kinase activity

GO:0000123 histone acetyltransferase complex

GO:0000151 ubiquitin ligase complex

GO:0000187 activation of MAPK activity

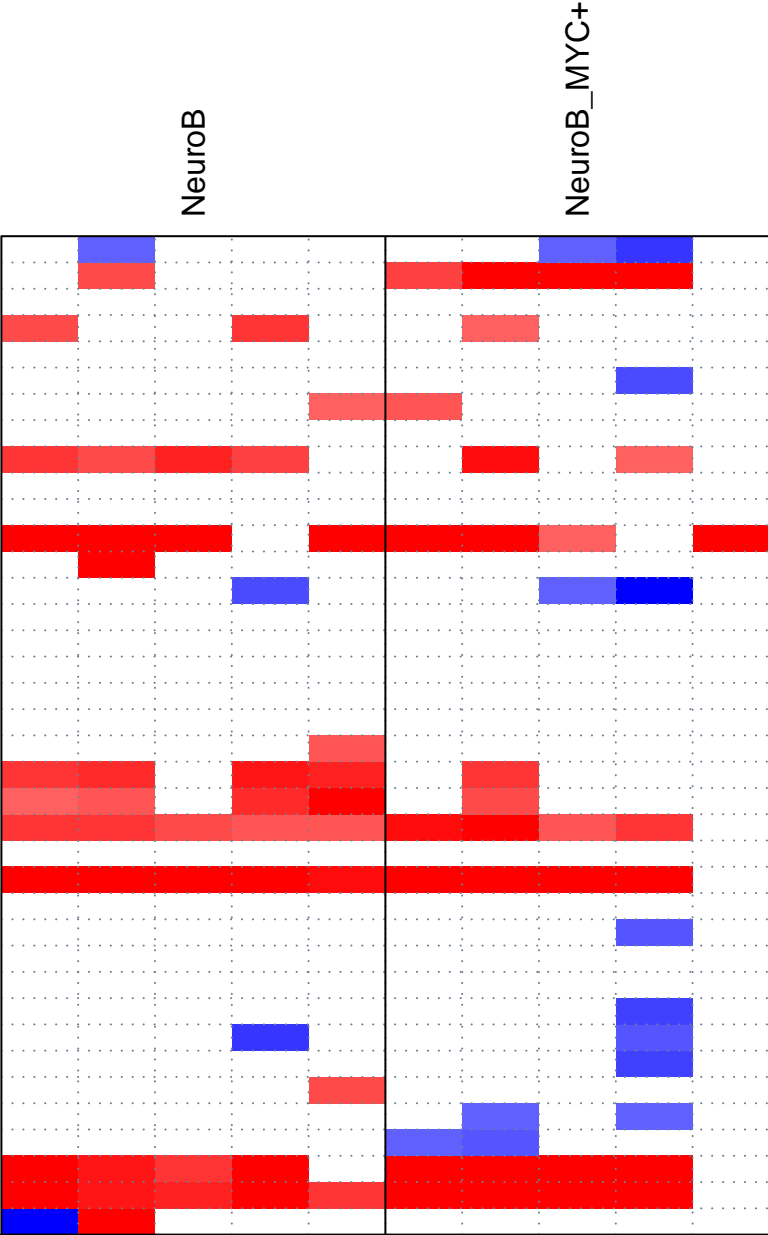
GO:0000226 microtubule cytoskeleton organization

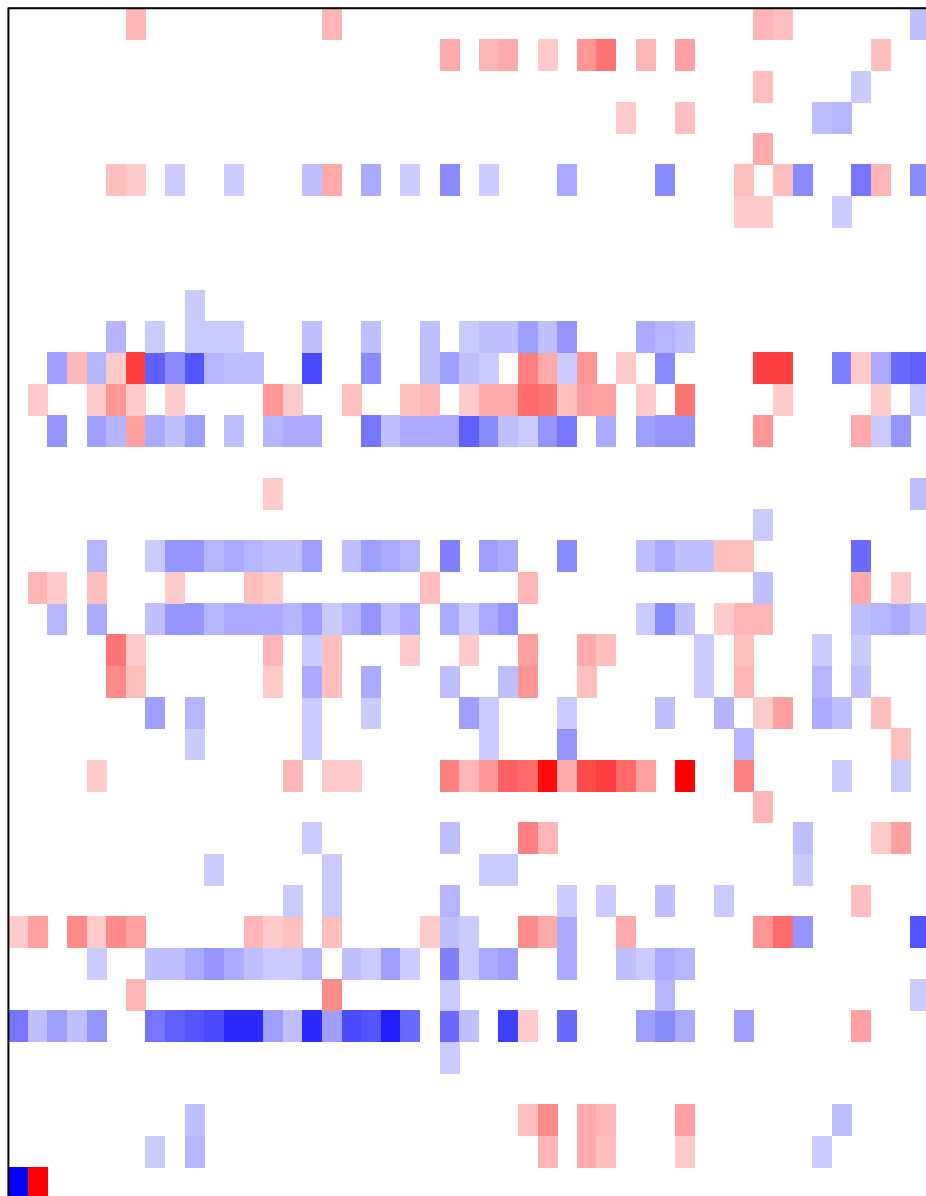
GO:0000228 nuclear chromosome

GO:0000271 polysaccharide biosynthetic process

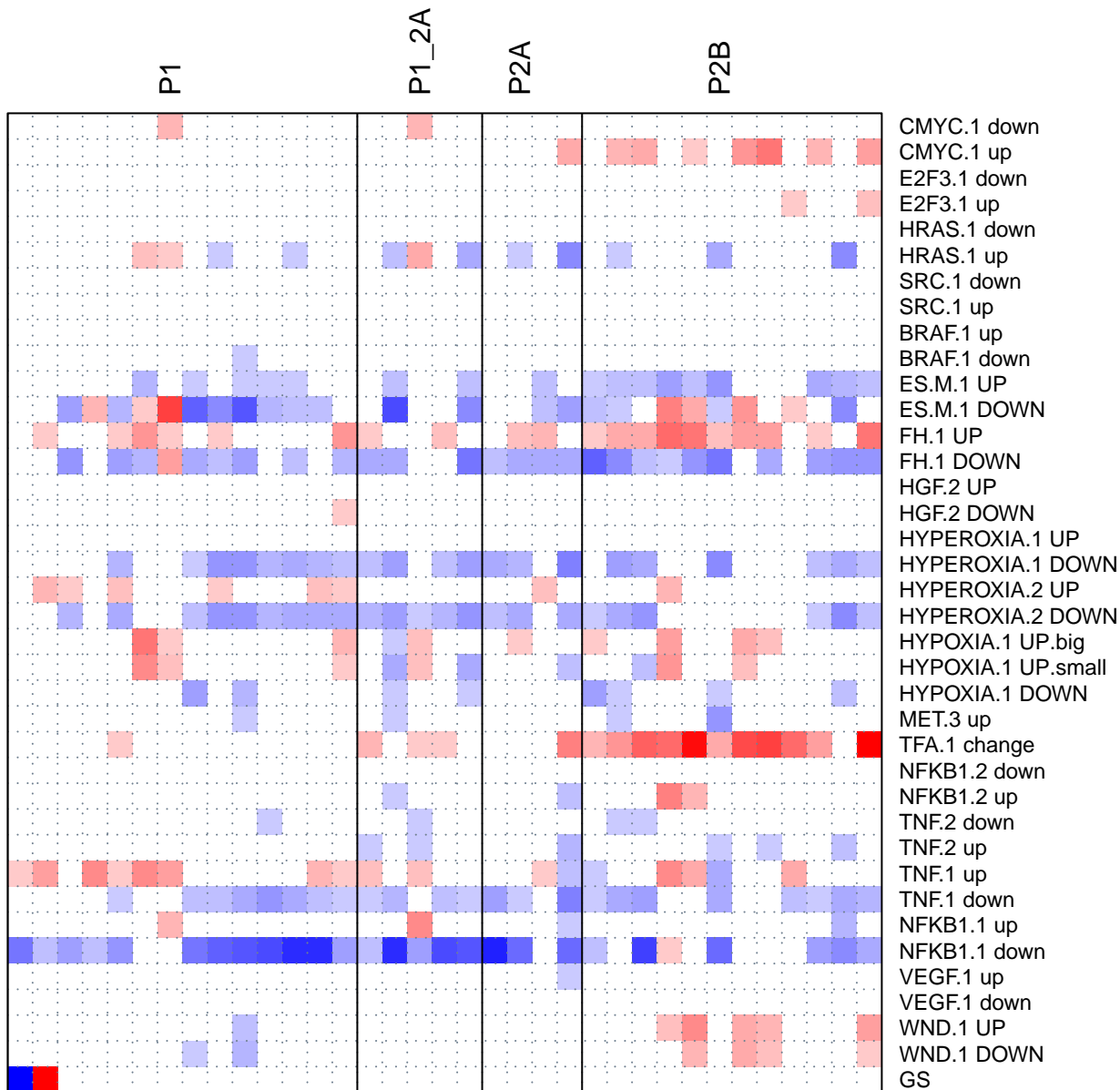
GO:0000302 response to reactive oxygen species

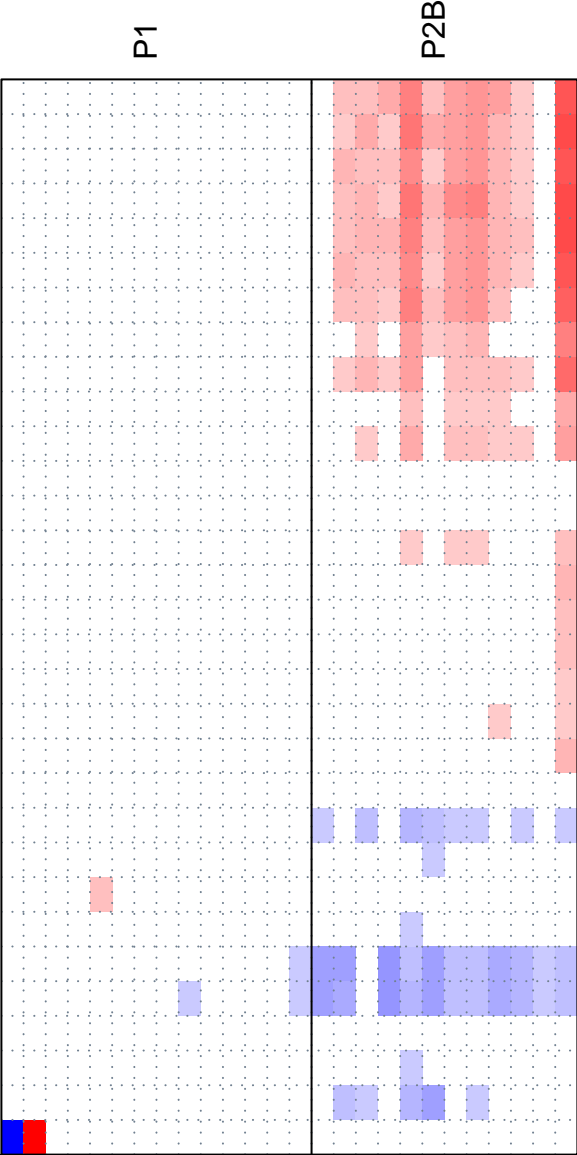
GS





CMYC.1 down
CMYC.1 up
E2F3.1 down
E2F3.1 up
HRAS.1 down
HRAS.1 up
SRC.1 down
SRC.1 up
BRAF.1 up
BRAF.1 down
ES.M.1 UP
ES.M.1 DOWN
FH.1 UP
FH.1 DOWN
HGF.2 UP
HGF.2 DOWN
HYPEROXIA.1 UP
HYPEROXIA.1 DOWN
HYPEROXIA.2 UP
HYPEROXIA.2 DOWN
HYPOXIA.1 UP.big
HYPOXIA.1 UP.small
HYPOXIA.1 DOWN
MET.3 up
TFA.1 change
NFKB1.2 down
NFKB1.2 up
TNF.2 down
TNF.2 up
TNF.1 up
TNF.1 down
NFKB1.1 up
NFKB1.1 down
VEGF.1 up
VEGF.1 down
WND.1 UP
WND.1 DOWN
GS





GO:0007059 chromosome segregation

GO:0005819 spindle

GO:0000779 condensed chromosome, centromeric region

GO:0000775 chromosome, centromeric region

GO:0000793 condensed chromosome

GO:0000777 condensed chromosome kinetochore

GO:0000776 kinetochore

GO:0000922 spindle pole

GO:0000226 microtubule cytoskeleton organization

GO:0010564 regulation of cell cycle process

GO:0007346 regulation of mitotic cell cycle

GO:0007088 regulation of mitosis

GO:0051783 regulation of nuclear division

GO:0051656 establishment of organelle localization

GO:0051321 meiotic cell cycle

GO:0007126 meiosis

GO:0051327 M phase of meiotic cell cycle

GO:0051640 organelle localization

GO:0048015 phosphoinositide-mediated signaling

GO:0000794 condensed nuclear chromosome

GO:0009416 response to light stimulus

GO:0048608 reproductive structure development

GO:0030216 keratinocyte differentiation

GO:0051047 positive regulation of secretion

GO:0043010 camera-type eye development

GO:0016327 apicolateral plasma membrane

GO:0043296 apical junction complex

GO:0021537 telencephalon development

GO:0001654 eye development

GO:0006720 isoprenoid metabolic process

GS