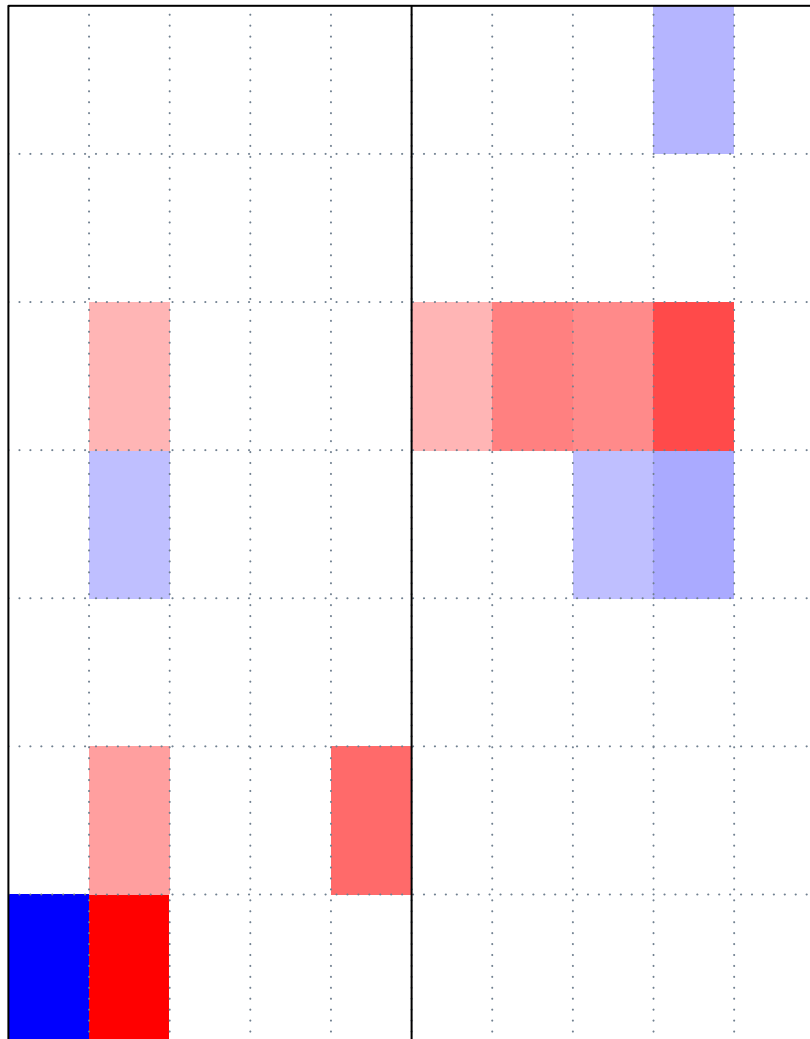


NeuroB

NeuroB_MYC+



ras UP – pmid: 16273092 NA

ras DN – pmid: 16273092 NA

myc UP – pmid: 16273092 NA

myc DN – pmid: 16273092 NA

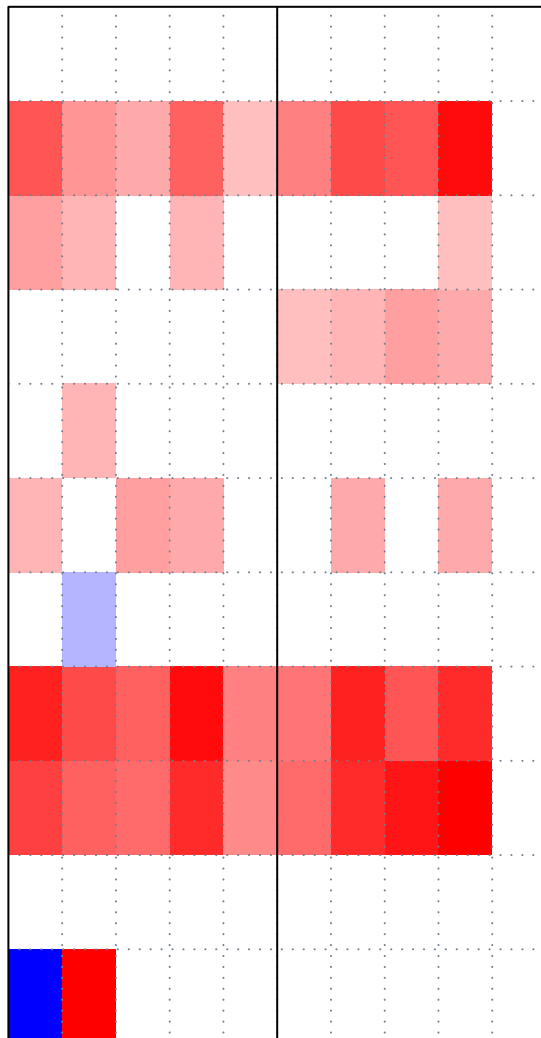
5p NA

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:0000139 Golgi membrane

GO:0000151 ubiquitin ligase complex

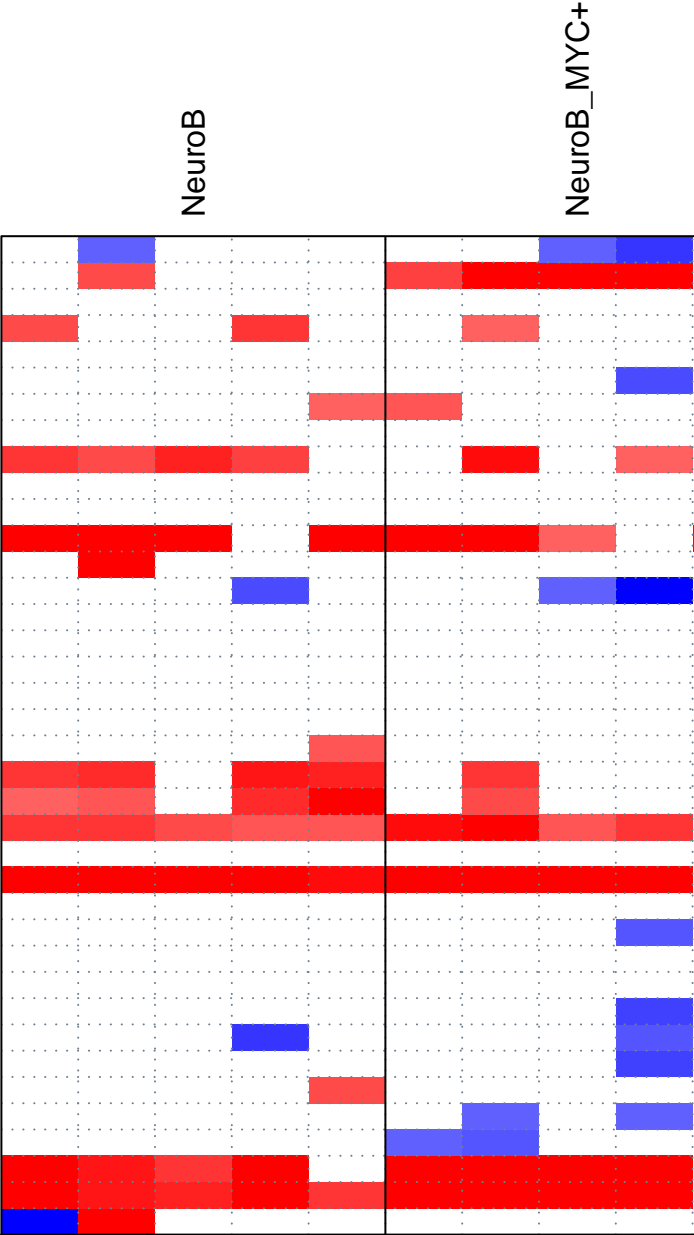
GO:0000187 activation of MAPK activity

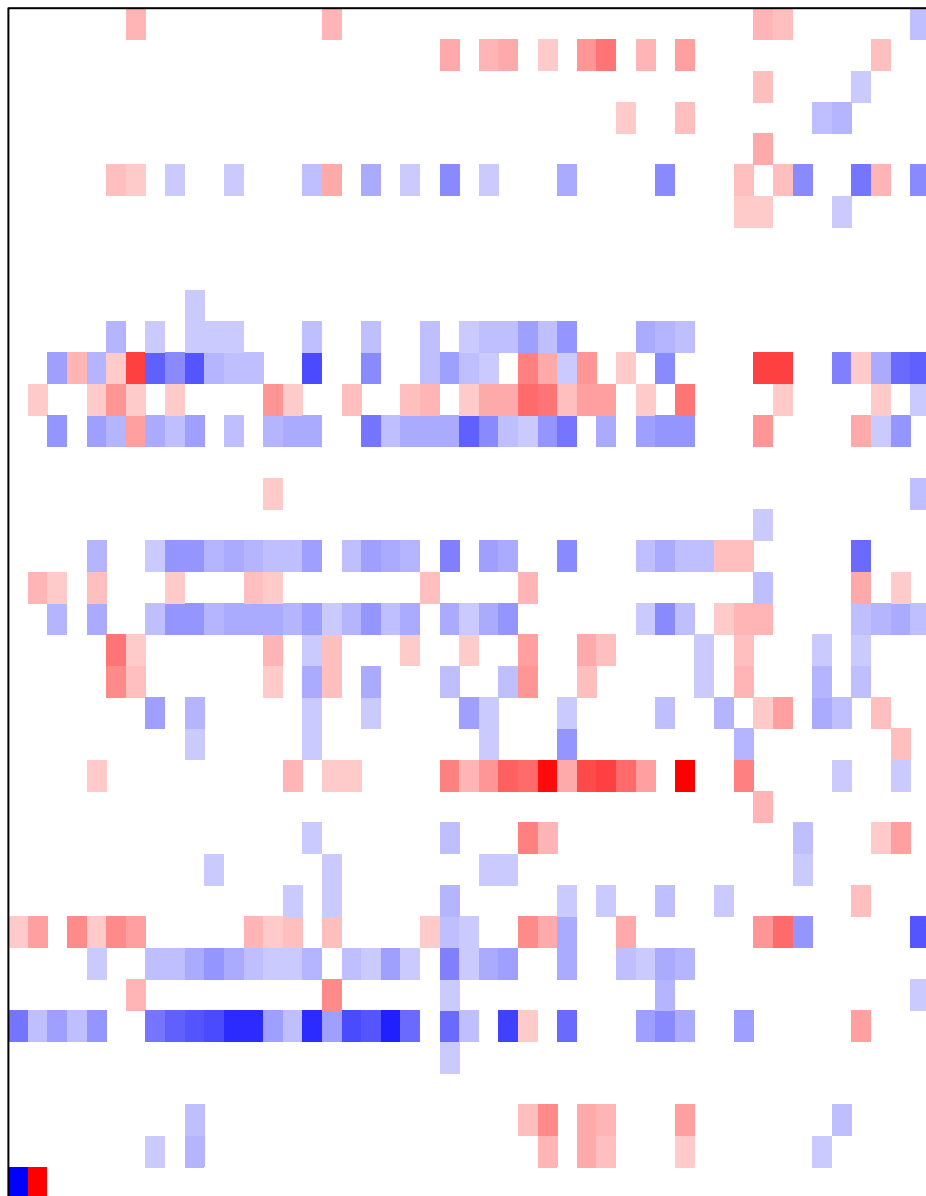
GO:0000226 microtubule cytoskeleton organization

GO:0000228 nuclear chromosome

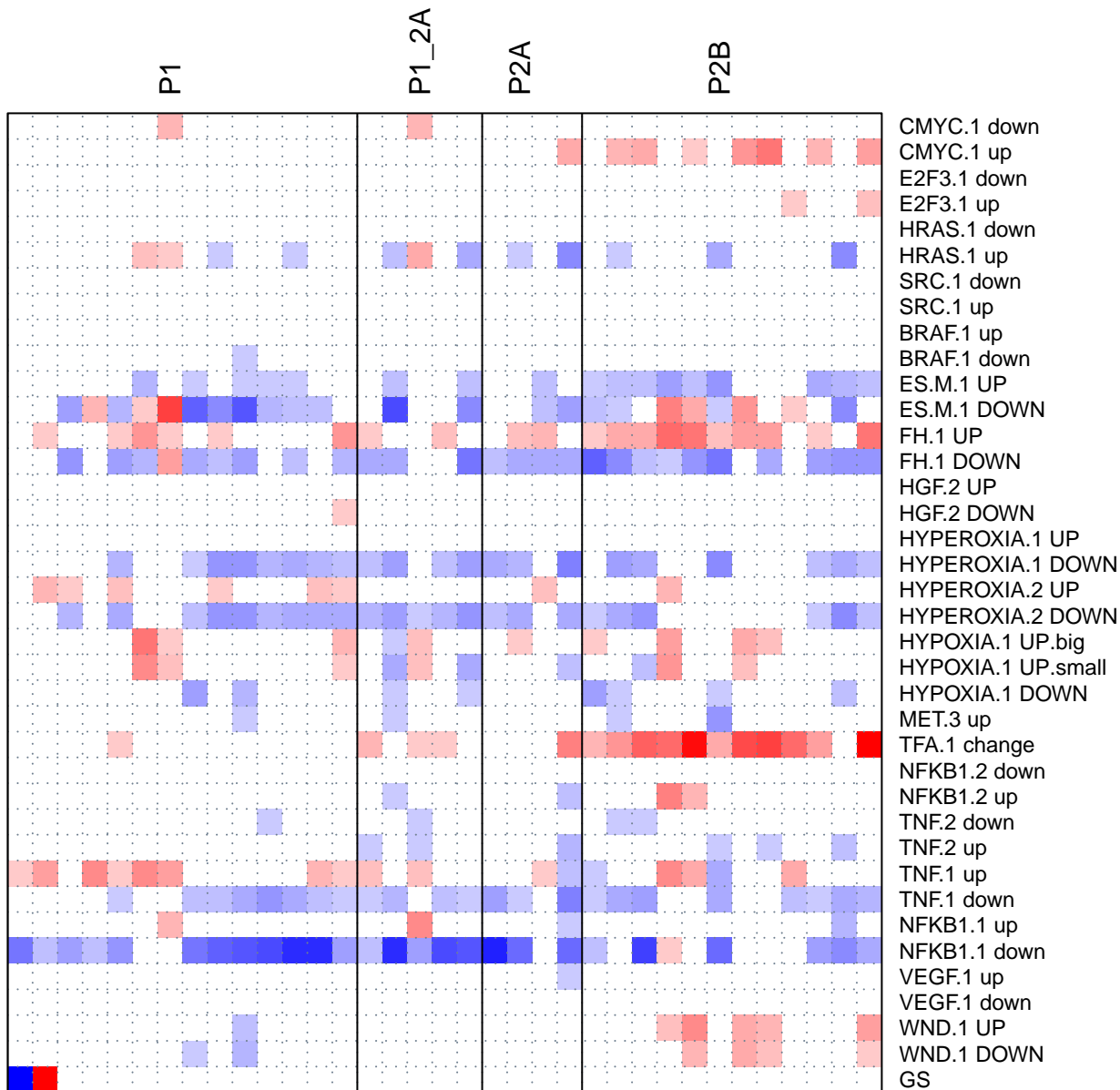
GO:0000271 polysaccharide biosynthetic process

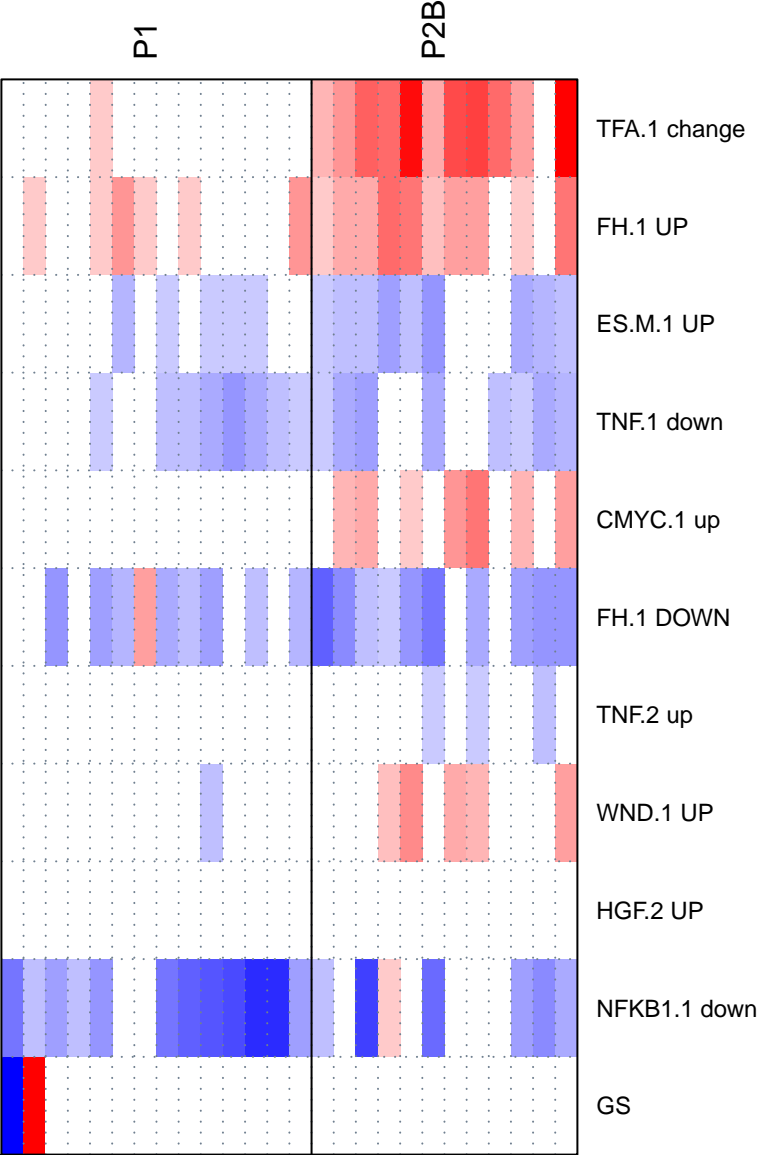
GS

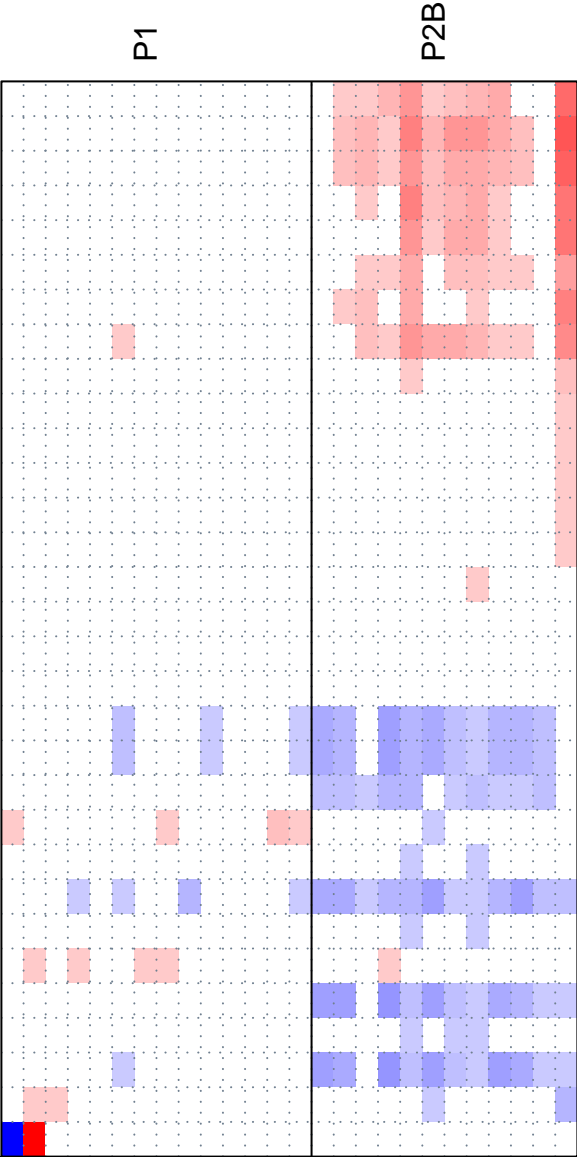




- CMYC.1 down
- CMYC.1 up
- E2F3.1 down
- E2F3.1 up
- HRAS.1 down
- HRAS.1 up
- SRC.1 down
- SRC.1 up
- BRAF.1 up
- BRAF.1 down
- ES.M.1 UP
- ES.M.1 DOWN
- FH.1 UP
- FH.1 DOWN
- HGF.2 UP
- HGF.2 DOWN
- HYPEROXIA.1 UP
- HYPEROXIA.1 DOWN
- HYPEROXIA.2 UP
- HYPEROXIA.2 DOWN
- HYPOXIA.1 UP.big
- HYPOXIA.1 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:0000775 chromosome, centromeric region

GO:0000793 condensed chromosome

GO:0005819 spindle

GO:0000776 kinetochore

GO:0007346 regulation of mitotic cell cycle

GO:0000226 microtubule cytoskeleton organization

GO:0000075 cell cycle checkpoint

GO:0010564 regulation of cell cycle process

GO:0007088 regulation of mitosis

GO:0051783 regulation of nuclear division

GO:0051321 meiotic cell cycle

GO:0007126 meiosis

GO:0051327 M phase of meiotic cell cycle

GO:0016763 transferase activity, transferring pentosyl groups

GO:0000079 regulation of cyclin-dependent protein kinase activity

GO:0032990 cell part morphogenesis

GO:0048592 eye morphogenesis

GO:0005923 tight junction

GO:0070160 occluding junction

GO:0001655 urogenital system development

GO:0030216 keratinocyte differentiation

GO:0043010 camera-type eye development

GO:0005911 cell-cell junction

GO:0007423 sensory organ development

GO:0051047 positive regulation of secretion

GO:0016327 apicolateral plasma membrane

GO:0001654 eye development

GO:0043296 apical junction complex

GO:0004553 hydrolase activity, hydrolyzing O-glycosyl compounds

GS