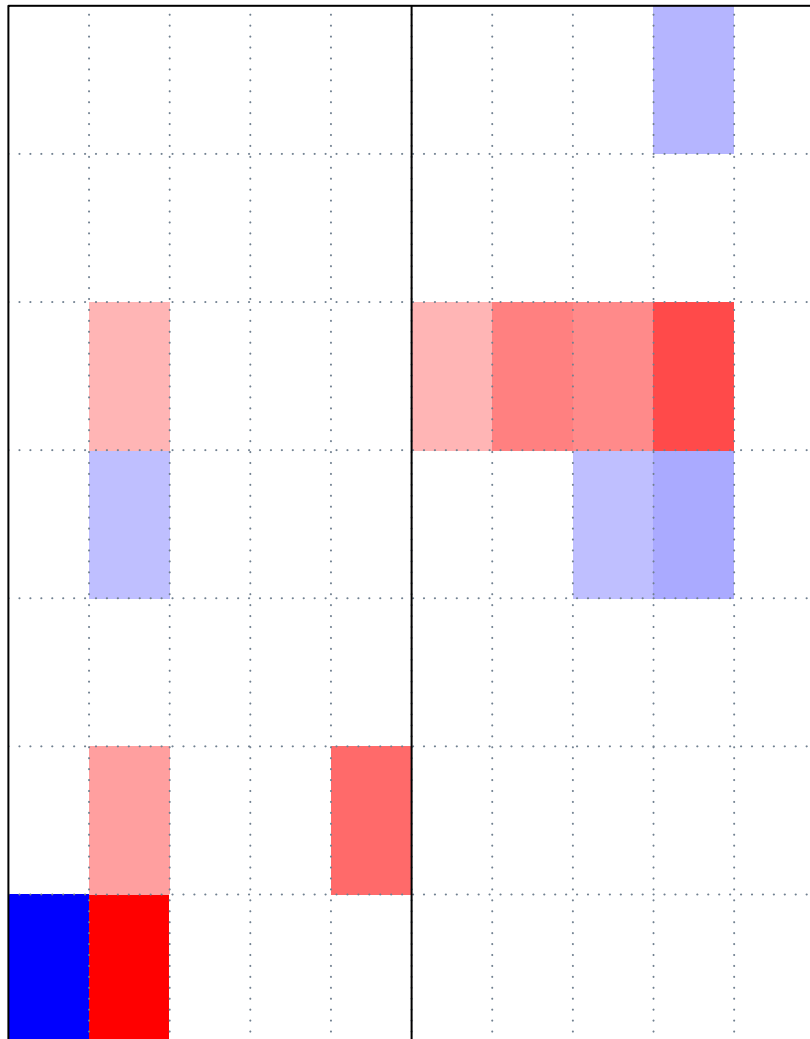


NeuroB

NeuroB_MYC+



ras UP – pmid: 16273092 NA

ras DN – pmid: 16273092 NA

myc UP – pmid: 16273092 NA

myc DN – pmid: 16273092 NA

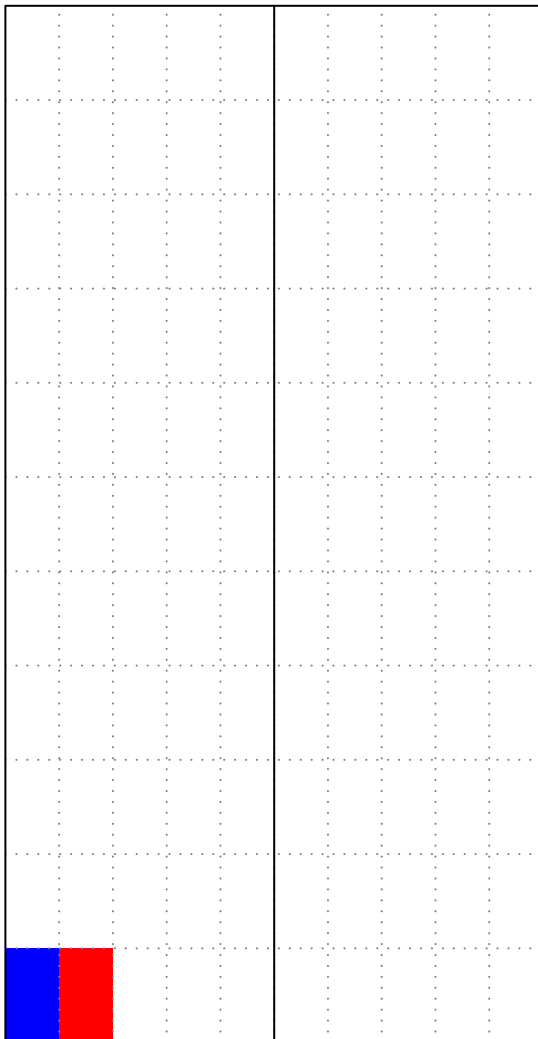
5p NA

5q NA

GS

NeuroB

NeuroB_MYC+



GO:0000002 mitochondrial genome maintenance

GO:0000018 regulation of DNA recombination

GO:0000022 mitotic spindle elongation

GO:0000027 ribosomal large subunit assembly and m

GO:0000030 mannosyltransferase activity

GO:0000049 tRNA binding

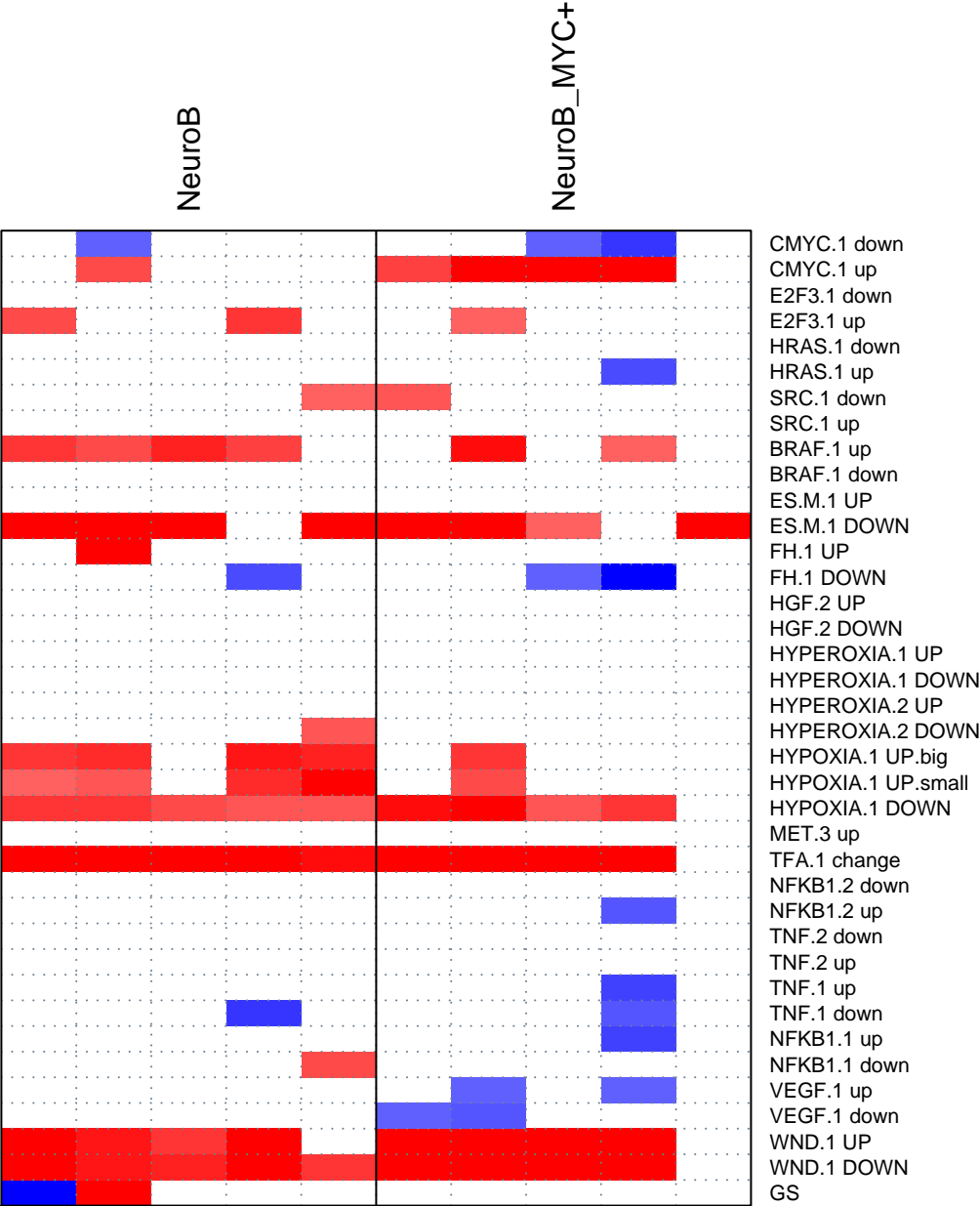
GO:0000051 urea cycle intermediate metabolic proces

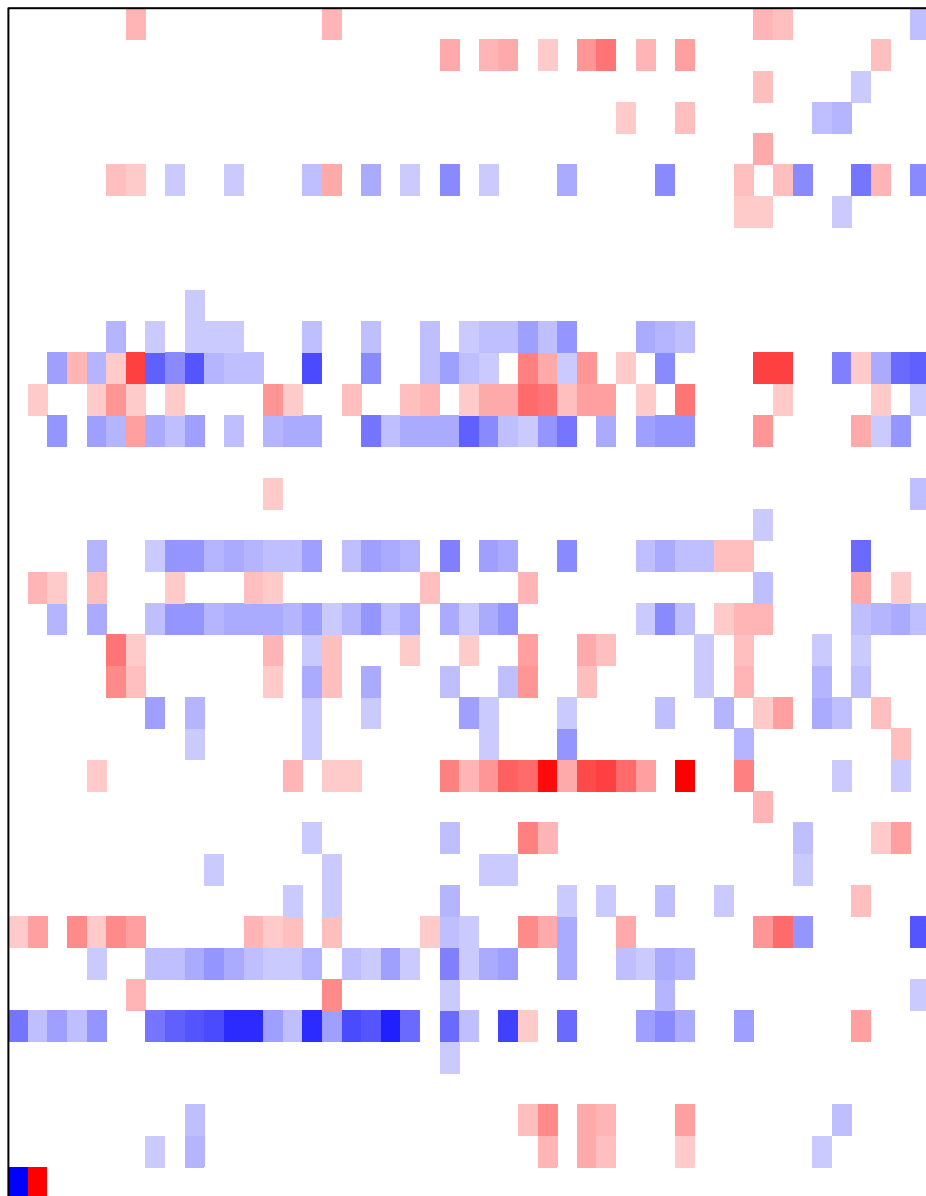
GO:0000059 protein import into nucleus, docking

GO:0000060 protein import into nucleus, translocation

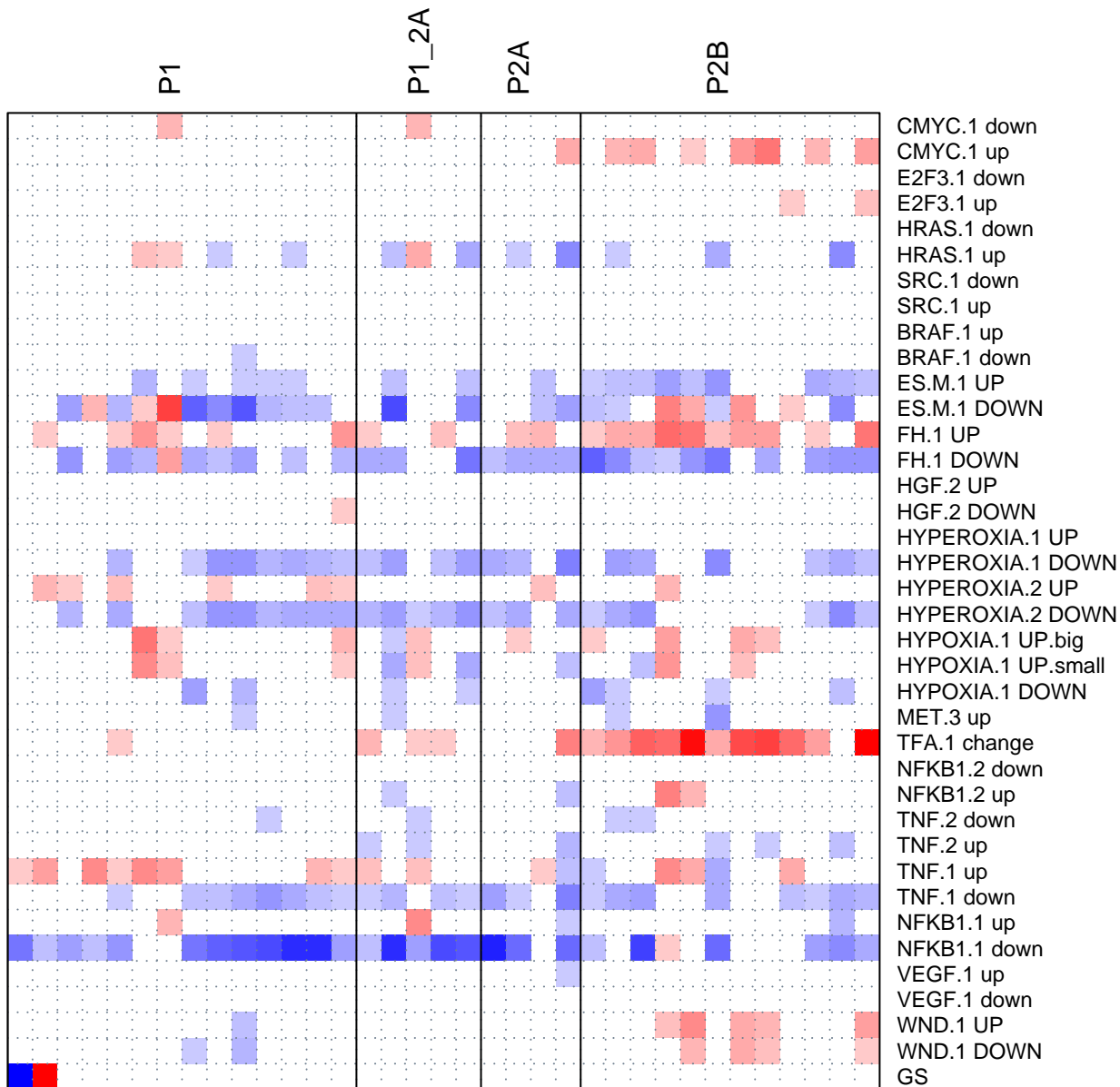
GO:0000062 acyl-CoA binding

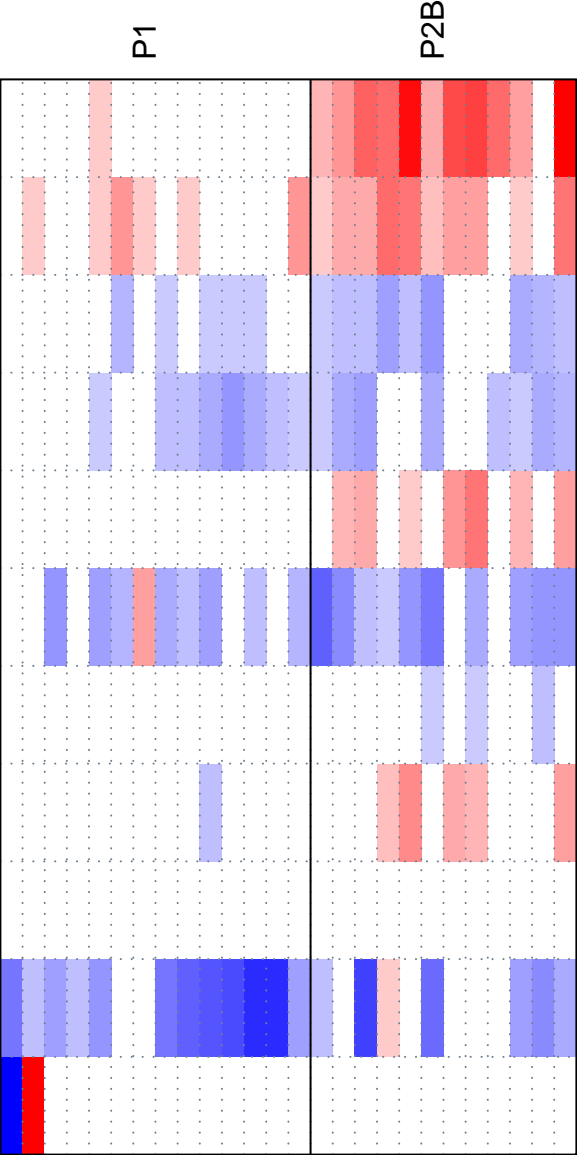
GS





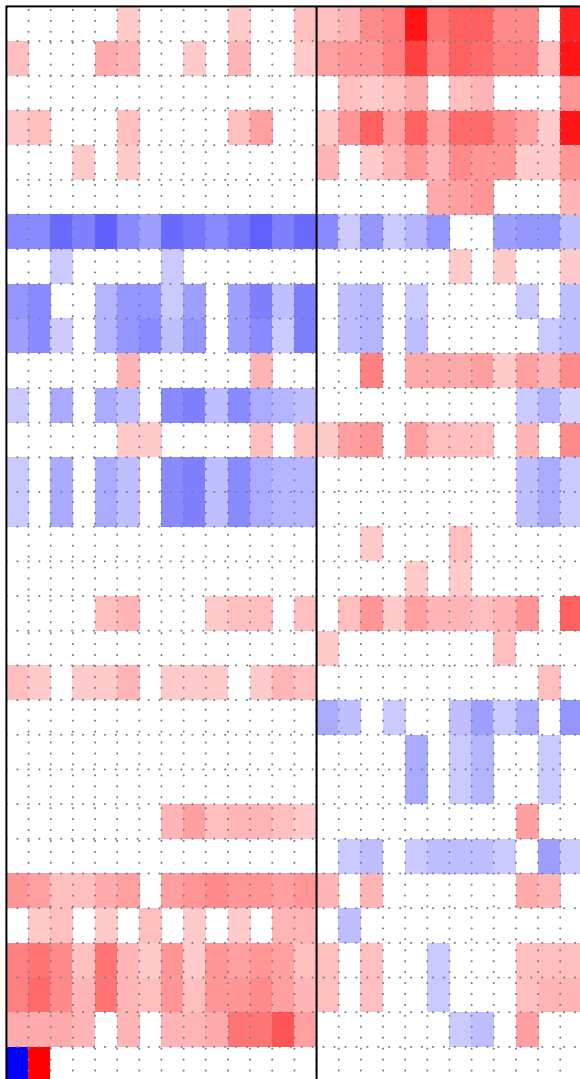
- CMYC.1 down
- CMYC.1 up
- E2F3.1 down
- E2F3.1 up
- HRAS.1 down
- HRAS.1 up
- SRC.1 down
- SRC.1 up
- BRAF.1 up
- BRAF.1 down
- ES.M.1 UP
- ES.M.1 DOWN
- FH.1 UP
- FH.1 DOWN
- HGF.2 UP
- HGF.2 DOWN
- HYPEROXIA.1 UP
- HYPEROXIA.1 DOWN
- HYPEROXIA.2 UP
- HYPEROXIA.2 DOWN
- HYPOXIA.1 UP.big
- HYPOXIA.1 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





P1

P2B



GO:0000776 kinetochore

GO:0007093 mitotic cell cycle checkpoint

GO:0007088 regulation of mitosis

GO:0006270 DNA replication initiation

GO:0000082 G1/S transition of mitotic cell cycle

GO:0006383 transcription from RNA polymerase III promoter

GO:0007173 epidermal growth factor receptor signaling pathway

GO:0016363 nuclear matrix

GO:0004114 3',5'-cyclic-nucleotide phosphodiesterase activity

GO:0004112 cyclic-nucleotide phosphodiesterase activity

GO:0003684 damaged DNA binding

GO:0051348 negative regulation of transferase activity

GO:0044450 microtubule organizing center part

GO:0006469 negative regulation of protein kinase activity

GO:0033673 negative regulation of kinase activity

GO:0035258 steroid hormone receptor binding

GO:0000079 regulation of cyclin-dependent protein kinase activity

GO:0031570 DNA integrity checkpoint

GO:0032313 regulation of Rab GTPase activity

GO:0009411 response to UV

GO:0005884 actin filament

GO:0030326 embryonic limb morphogenesis

GO:0035113 embryonic appendage morphogenesis

GO:0030286 dynein complex

GO:0030855 epithelial cell differentiation

GO:0019861 flagellum

GO:0005548 phospholipid transporter activity

GO:0016540 protein autoprocessing

GO:0046777 protein amino acid autophosphorylation

GO:0001764 neuron migration

GS