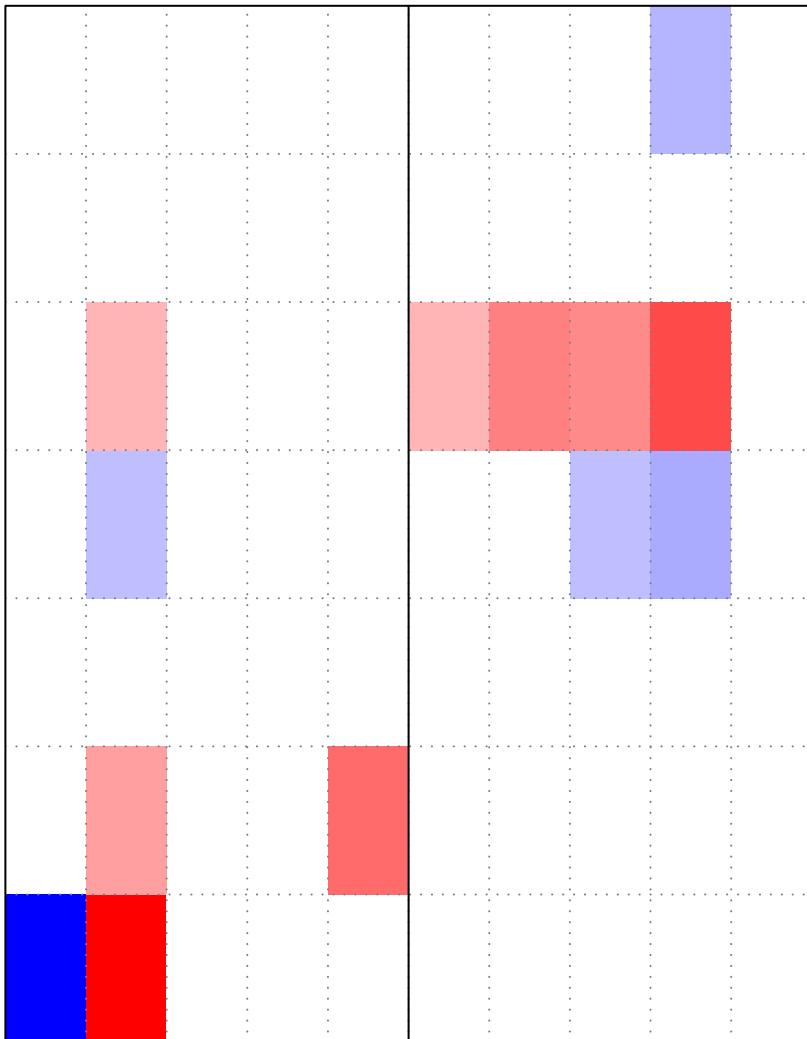


NeuroB

NeuroB_MYC+



ras UP – pmid: 16273092 NA

ras DN – pmid: 16273092 NA

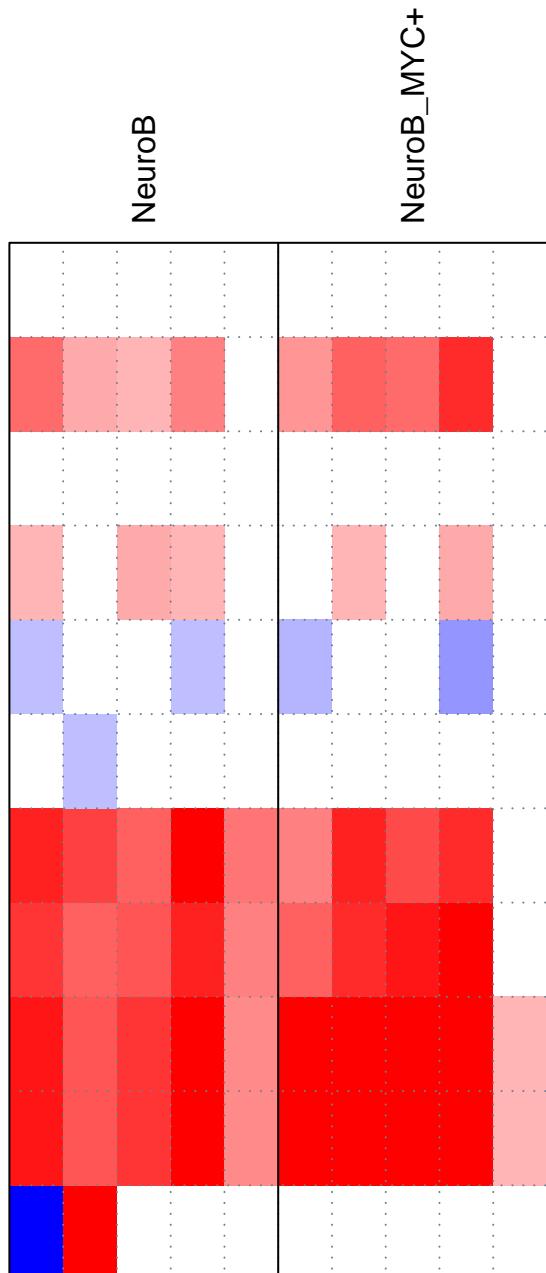
myc UP – pmid: 16273092 NA

myc DN – pmid: 16273092 NA

5p NA

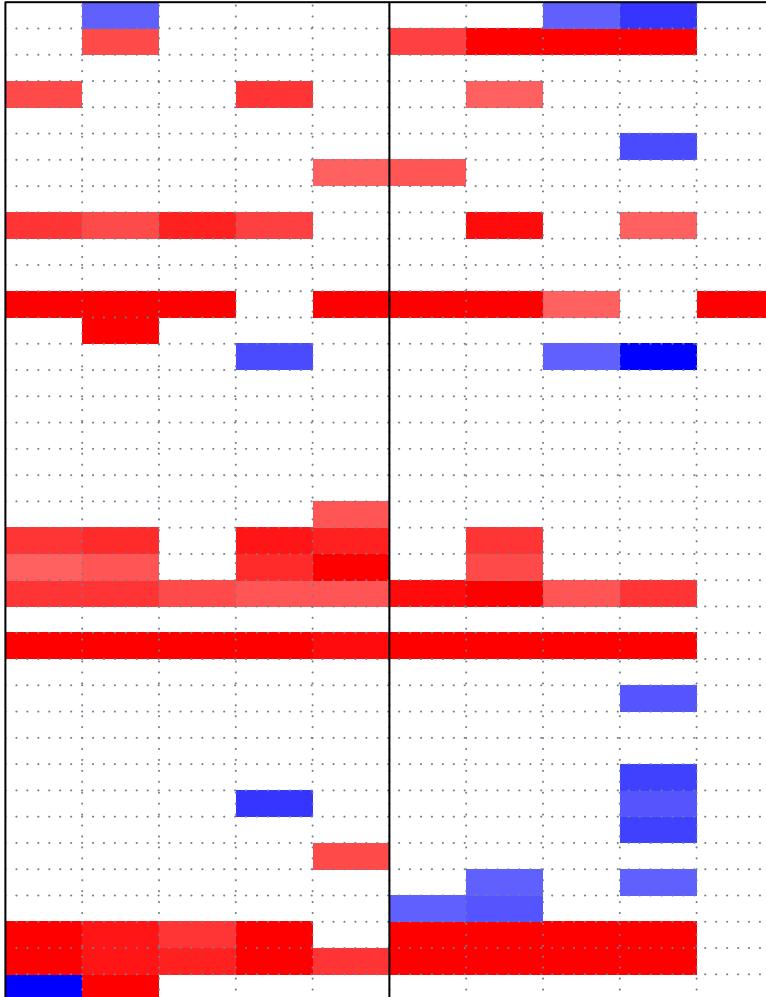
5q NA

GS



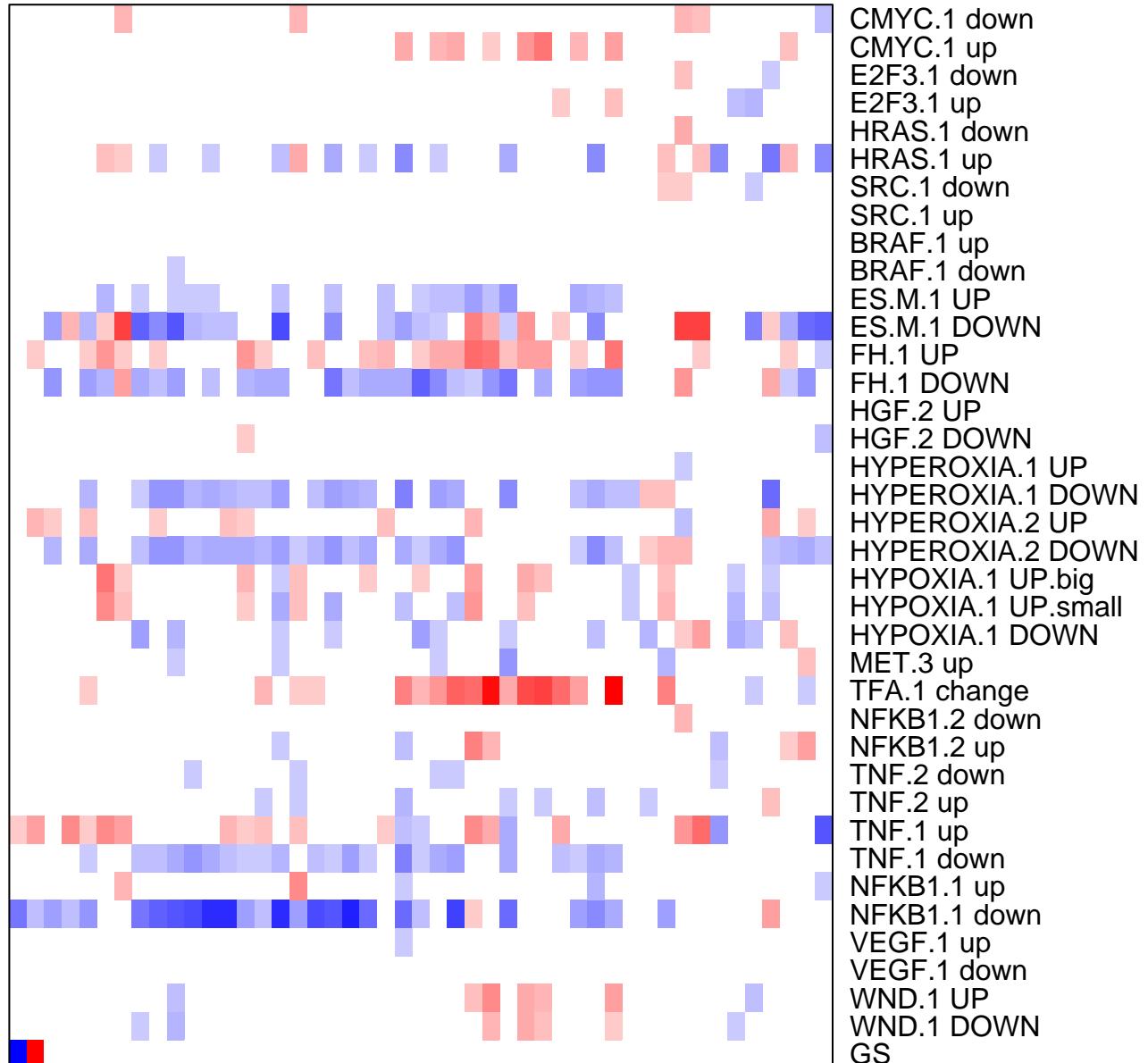
- GO:0000041 transition metal ion transport
- GO:0000075 cell cycle checkpoint
- GO:0000122 negative regulation of transcription from DNA template
- GO:0000151 ubiquitin ligase complex
- GO:0000165 MAPKKK cascade
- GO:0000187 activation of MAPK activity
- GO:0000226 microtubule cytoskeleton organization and maintenance
- GO:0000228 nuclear chromosome
- GO:0000375 RNA splicing, via transesterification reaction
- GO:0000377 RNA splicing, via transesterification reaction
- GS

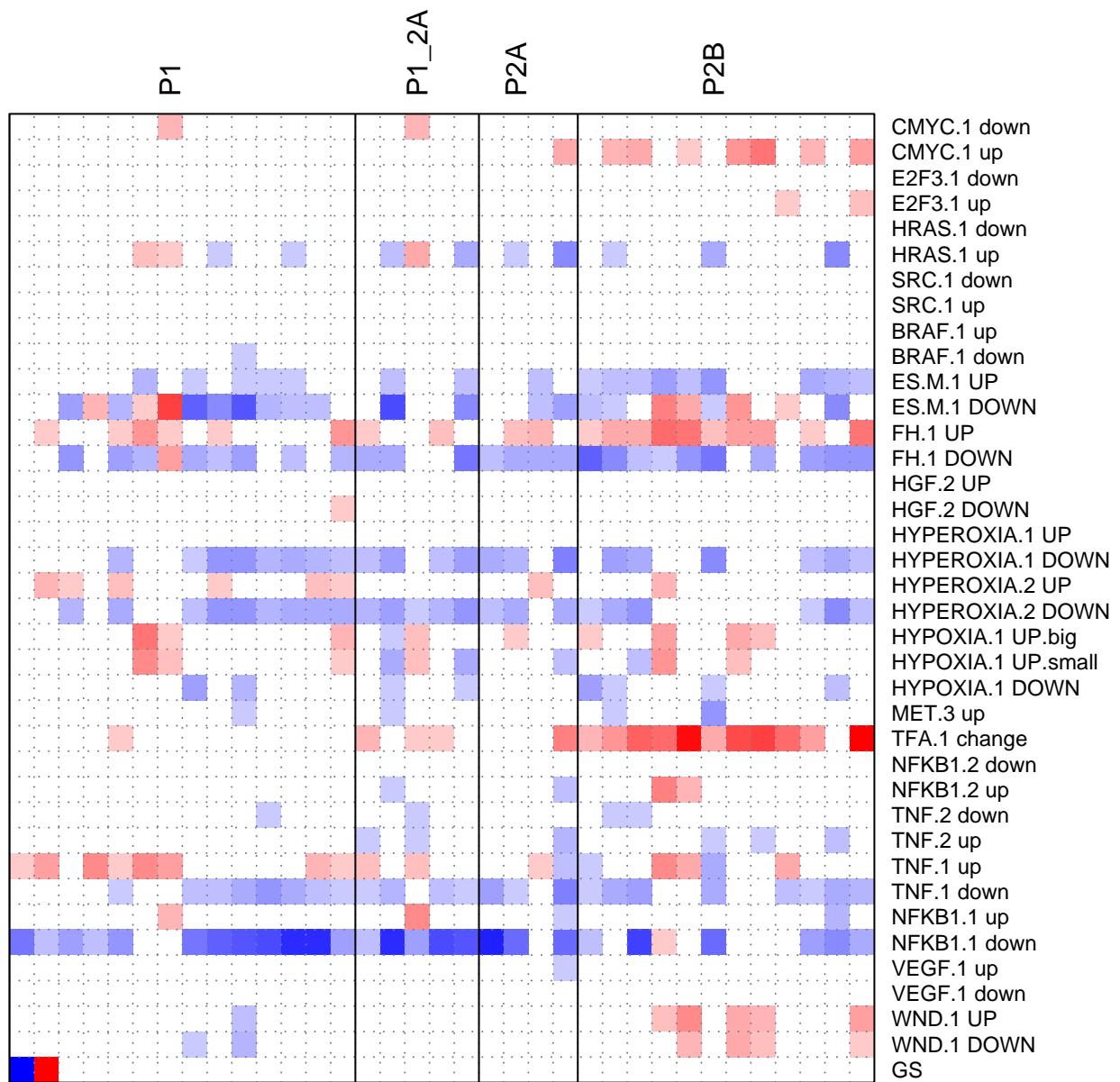
NeuroB



NeuroB_MYC+

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

TFA.1 change

FH.1 UP

ES.M.1 UP

TNF.1 down

CMYC.1 up

FH.1 DOWN

TNF.2 up

WND.1 UP

HGF.2 UP

NFKB1.1 down

GS

