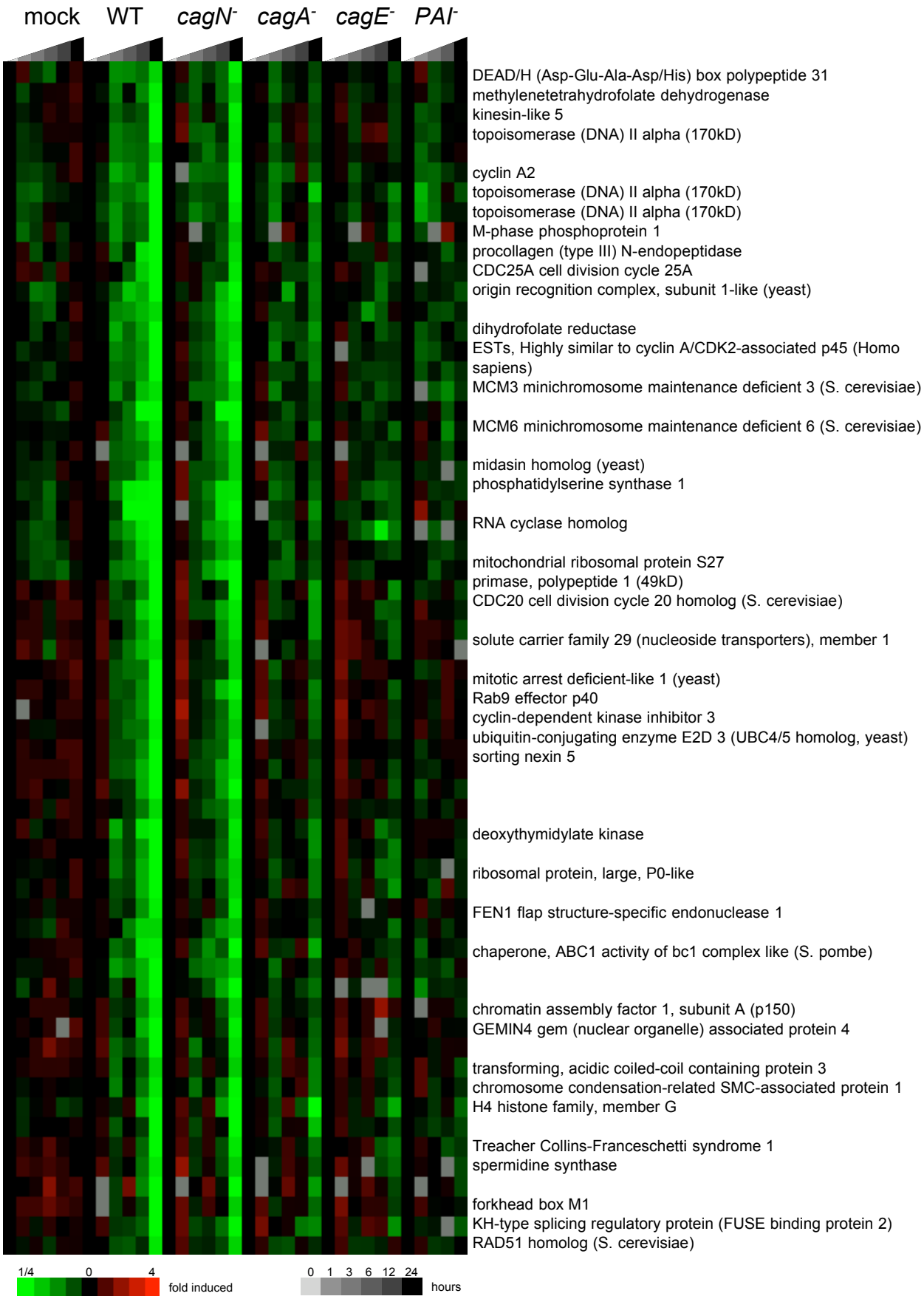
mock
WT
cagN-
cagA-
cagE-
PAI-
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31
methylenetetrahydrofolate dehydrogenase
kinesin-like 5
topoisomerase (DNA) II alpha (170kD)
cyclin A2
topoisomerase (DNA) II alpha (170kD)
topoisomerase (DNA) II alpha (170kD)
M-phase phosphoprotein 1
procollagen (type III) N-endopeptidase
CDC25A cell division cycle 25A
origin recognition complex, subunit 1-like (yeast)
dihydrofolate reductase
ESTs, Highly similar to cyclin A/CDK2-associated p45 (Homo sapiens)
MCM3 minichromosome maintenance deficient 3 (S. cerevisiae)
MCM6 minichromosome maintenance deficient 6 (S. cerevisiae)
midasin homolog (yeast)
phosphatidylserine synthase 1
RNA cyclase homolog
mitochondrial ribosomal protein S27
primase, polypeptide 1 (49kD)
CDC20 cell division cycle 20 homolog (S. cerevisiae)
solute carrier family 29 (nucleoside transporters), member 1
mitotic arrest deficient-like 1 (yeast)
Rab9 effector p40
cyclin-dependent kinase inhibitor 3
ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)
sorting nexin 5
deoxythymidylate kinase
ribosomal protein, large, P0-like
FEN1 flap structure-specific endonuclease 1
chaperone, ABC1 activity of bc1 complex like (S. pombe)
chromatin assembly factor 1, subunit A (p150)
GEMIN4 gem (nuclear organelle) associated protein 4
transforming, acidic coiled-coil containing protein 3
chromosome condensation-related SMC-associated protein 1
H4 histone family, member G
Treacher Collins-Franceschetti syndrome 1
spermidine synthase
forkhead box M1
KH-type splicing regulatory protein (FUSE binding protein 2)
RAD51 homolog (S. cerevisiae)
1/4
0
4
fold induced
0 1 3 6 12 24
hours