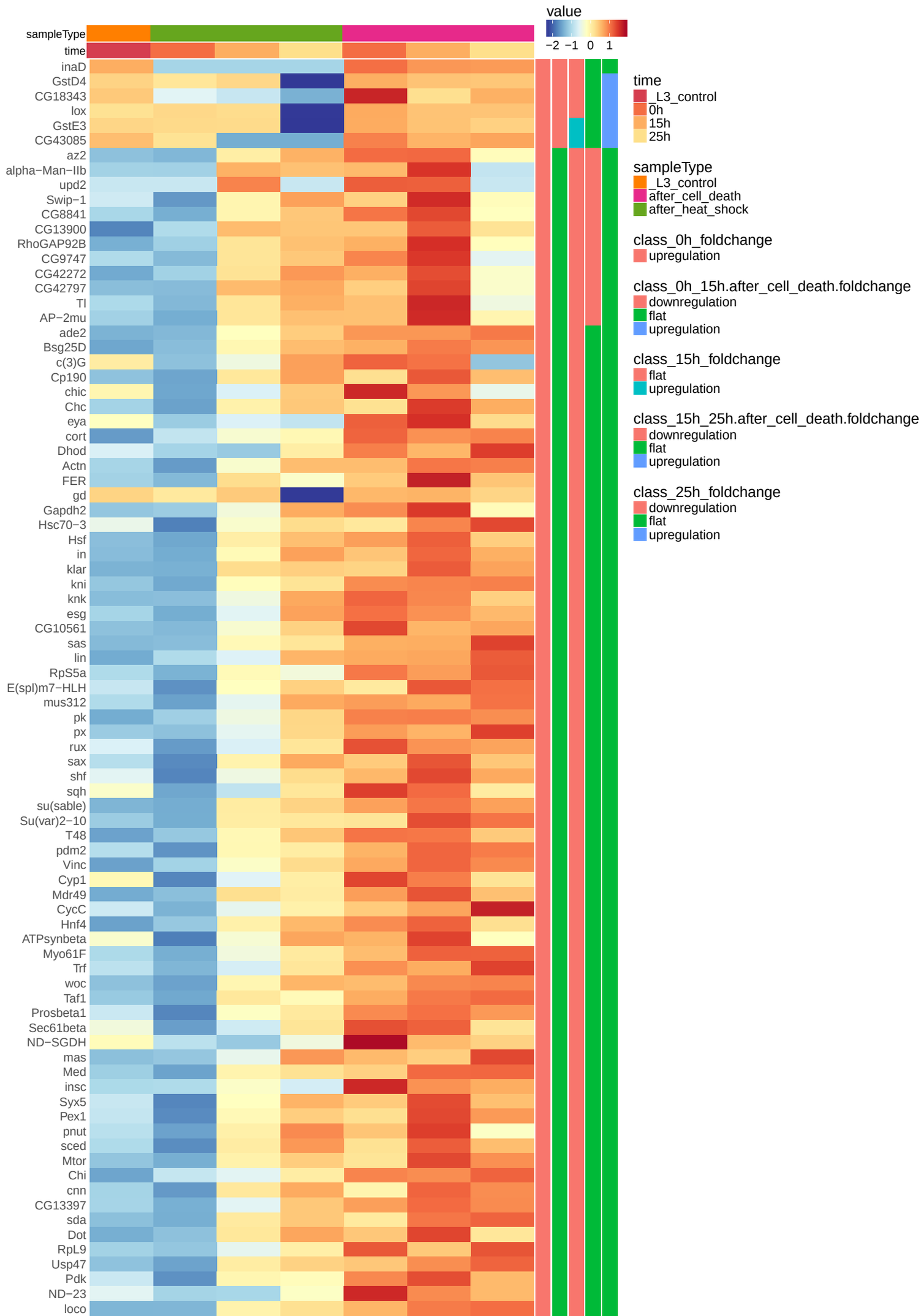
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Tl
AP−2mu
ade2
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c(3)G
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Chc
eya
cort
Dhod
Actn
FER
gd
Gapdh2
Hsc70−3
Hsf
in
klar
kni
knk
esg
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sas
lin
RpS5a
E(spl)m7−HLH
mus312
pk
px
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sax
shf
sqh
su(sable)
Su(var)2−10
T48
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Vinc
Cyp1
Mdr49
CycC
Hnf4
ATPsynbeta
Myo61F
Trf
woc
Taf1
Prosbeta1
Sec61beta
ND−SGDH
mas
Med
insc
Syx5
Pex1
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Mtor
Chi
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Dot
RpL9
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Pdk
ND−23
loco
time
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15h
25h
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after_heat_shock
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upregulation
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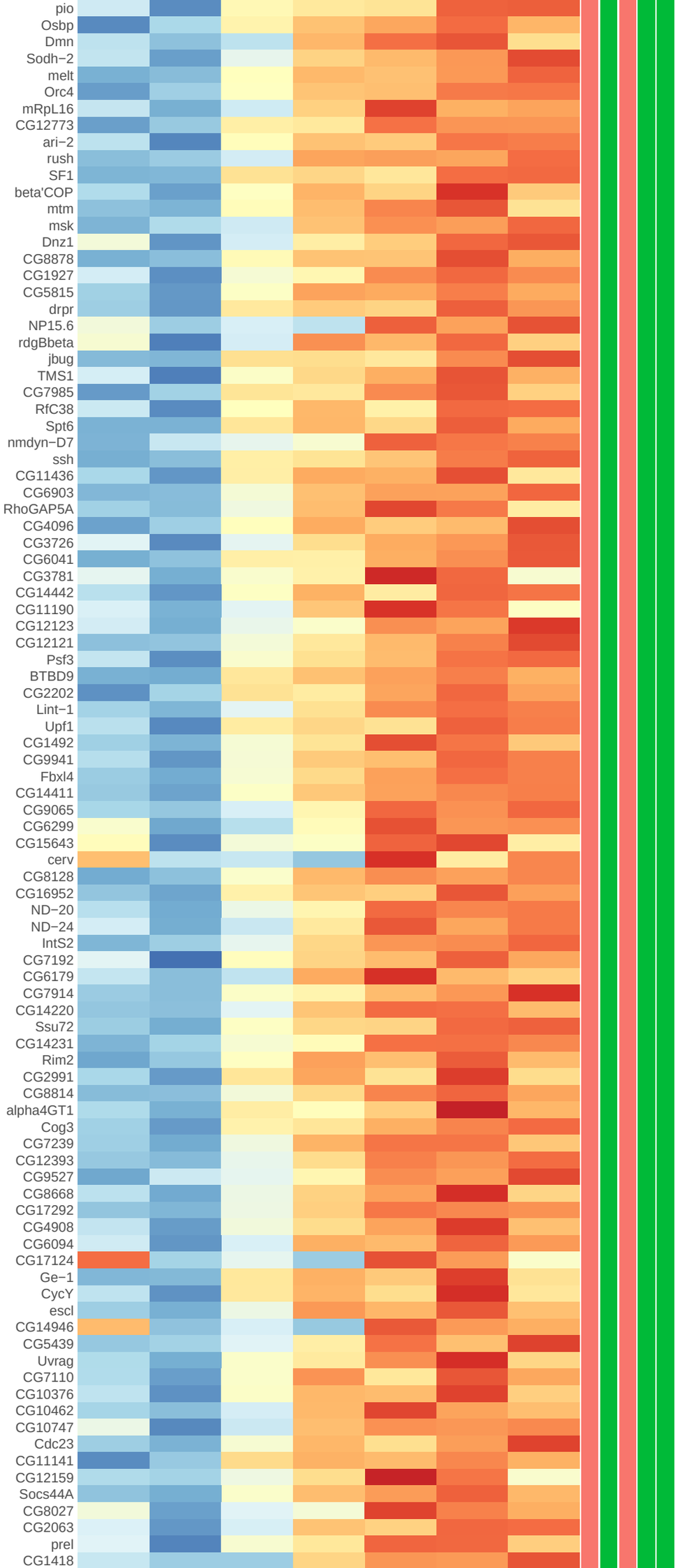
downregulation
flat
upregulation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Spt6
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CG2202
Lint−1
Upf1
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CycY
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Uvrag
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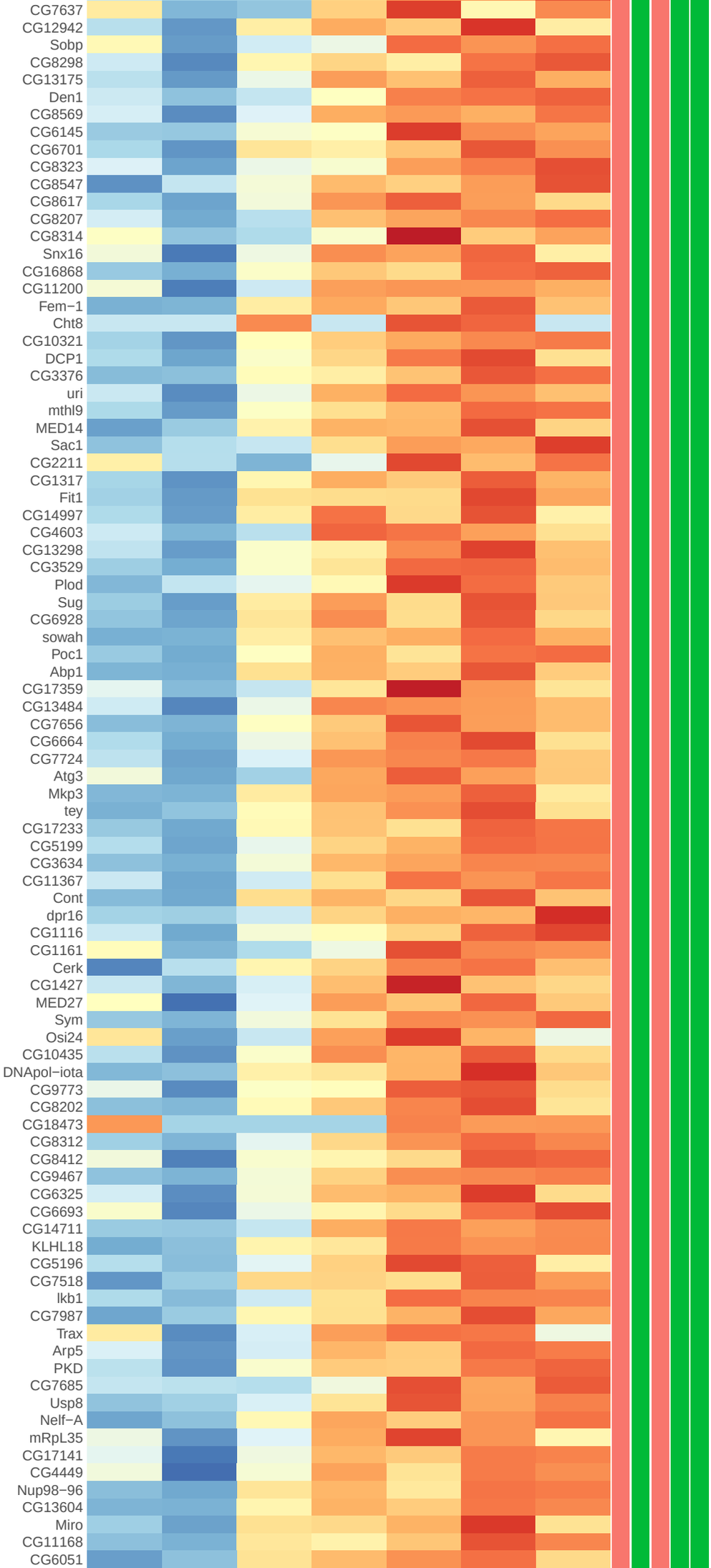
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MED14
Sac1
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Plod
Sug
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Poc1
Abp1
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CG13484
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Mkp3
tey
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Cont
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Cerk
CG1427
MED27
Sym
Osi24
CG10435
DNApol−iota
CG9773
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CG14711
KLHL18
CG5196
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Ikb1
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Trax
Arp5
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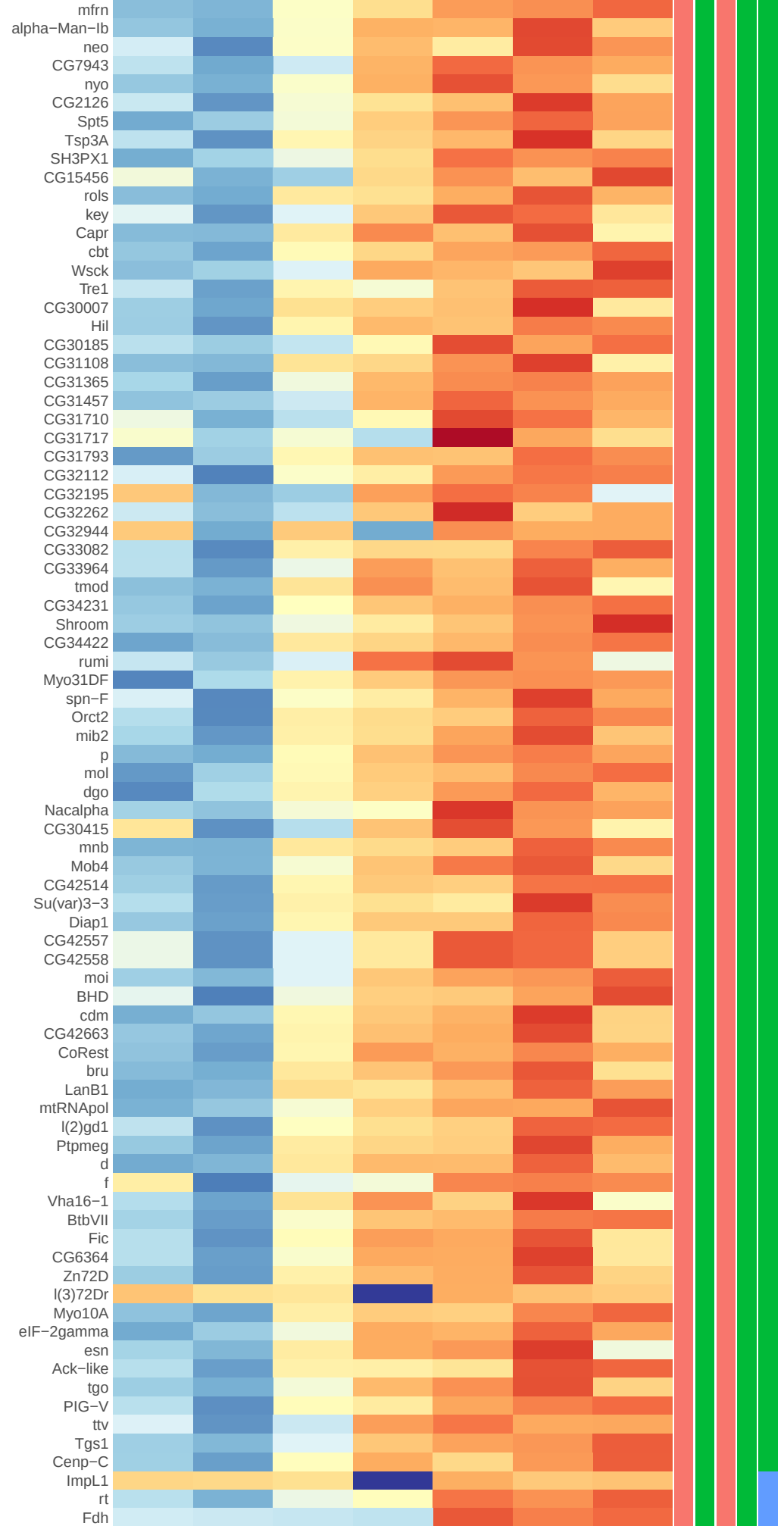
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cbt
Wsck
Tre1
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CG30185
CG31108
CG31365
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CG31710
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CG31793
CG32112
CG32195
CG32262
CG32944
CG33082
CG33964
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CG34231
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CG34422
rumi
Myo31DF
spn−F
Orct2
mib2
p
mol
dgo
Nacalpha
CG30415
mnb
Mob4
CG42514
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Diap1
CG42557
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BHD
cdm
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Ptpmeg
d
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Vha16−1
BtbVII
Fic
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Zn72D
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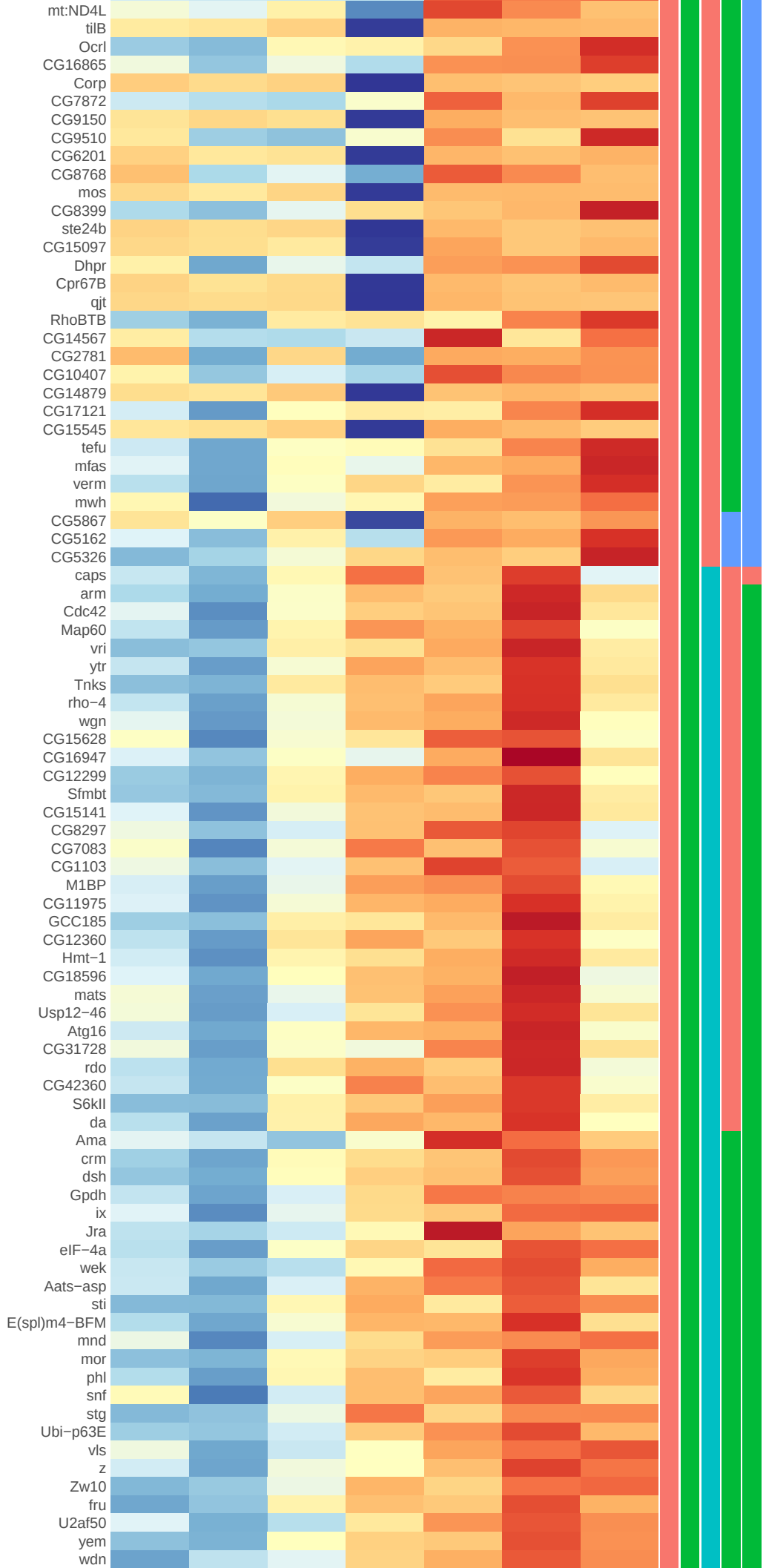
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dsh
Gpdh
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Jra
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wek
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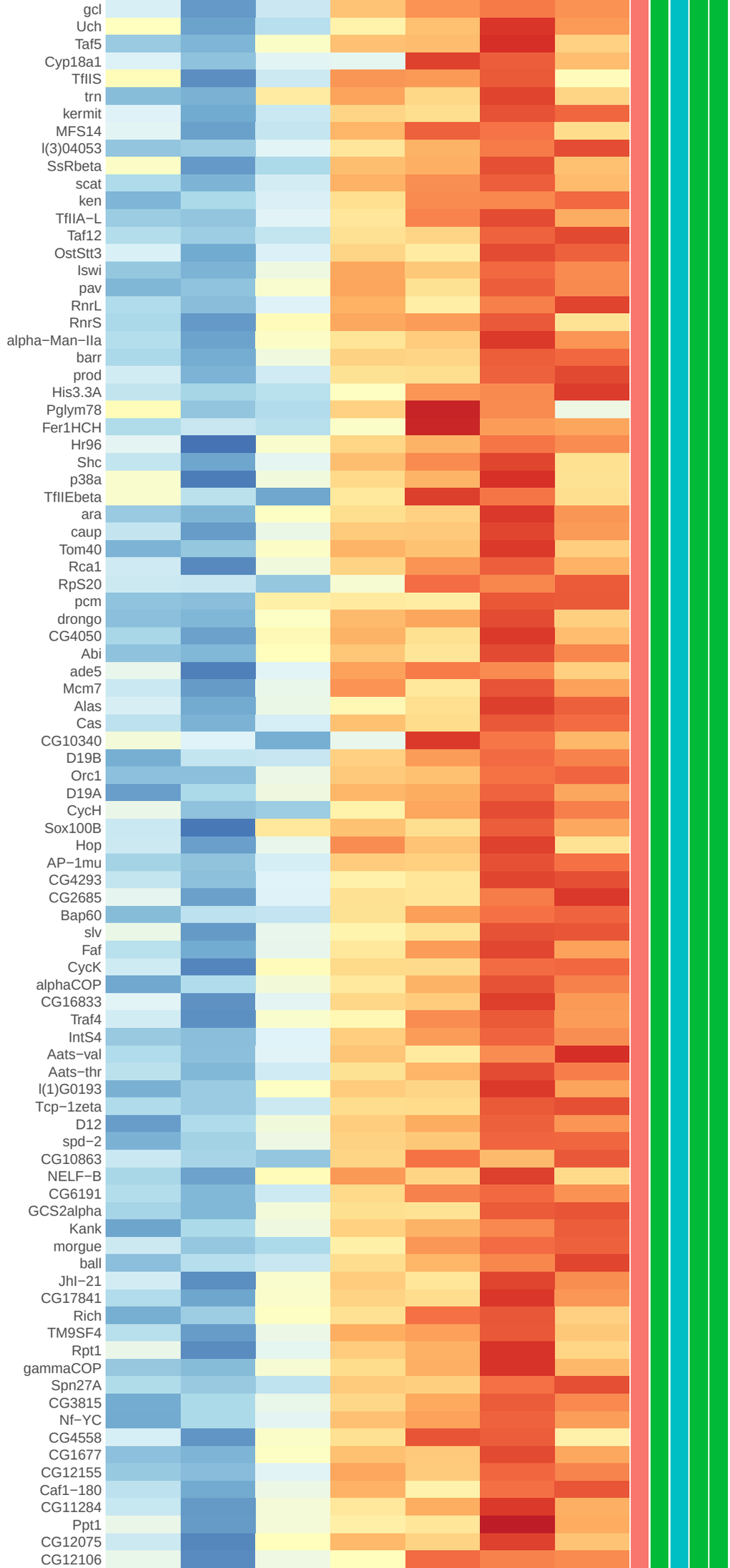
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Taf12
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RnrL
RnrS
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barr
prod
His3.3A
Pglym78
Fer1HCH
Hr96
Shc
p38a
TfIIEbeta
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caup
Tom40
Rca1
RpS20
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drongo
CG4050
Abi
ade5
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Alas
Cas
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Orc1
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CycH
Sox100B
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Rpt1
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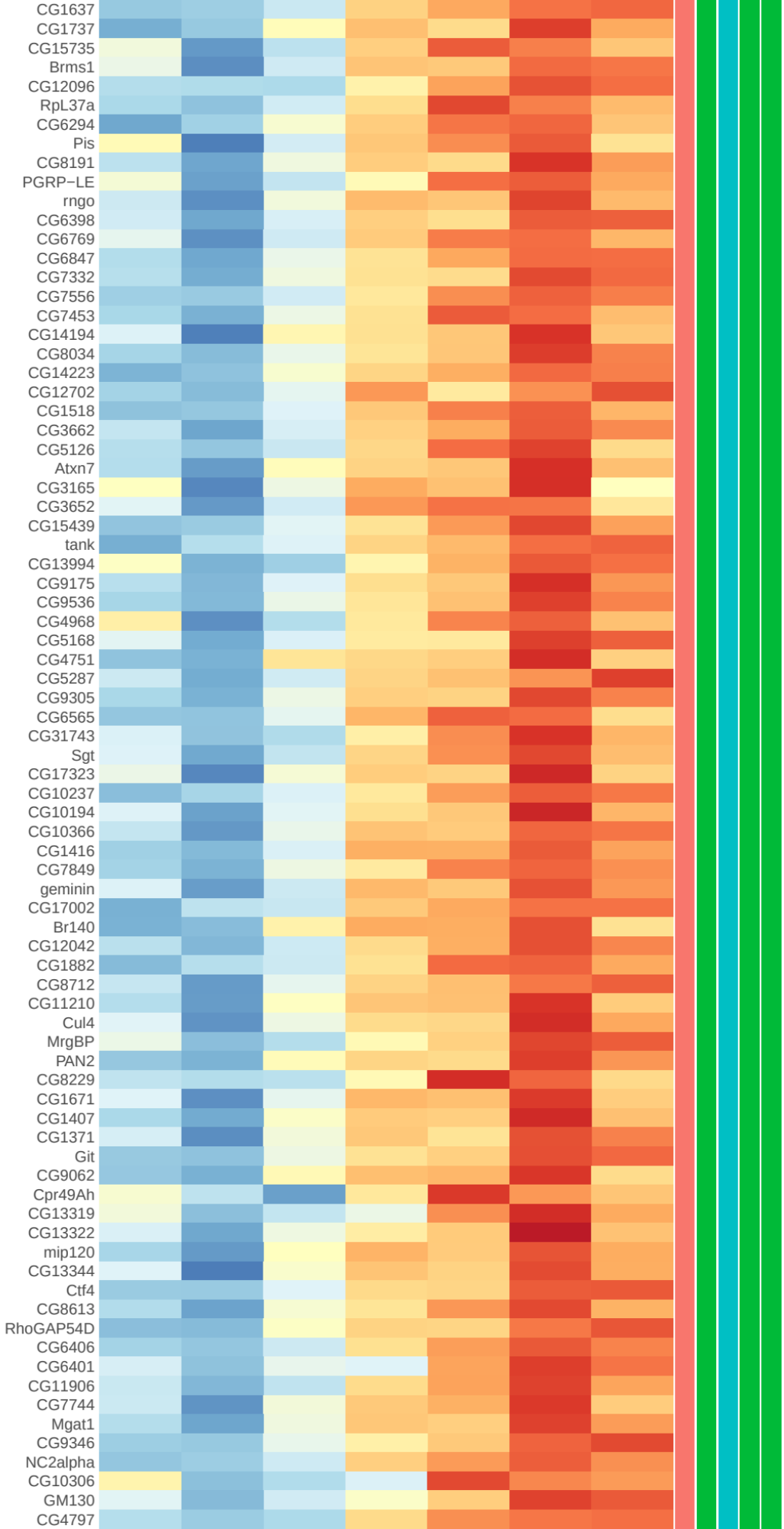
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CG3662
CG5126
Atxn7
CG3165
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CG12042
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PAN2
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CG1407
CG1371
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mip120
CG13344
Ctf4
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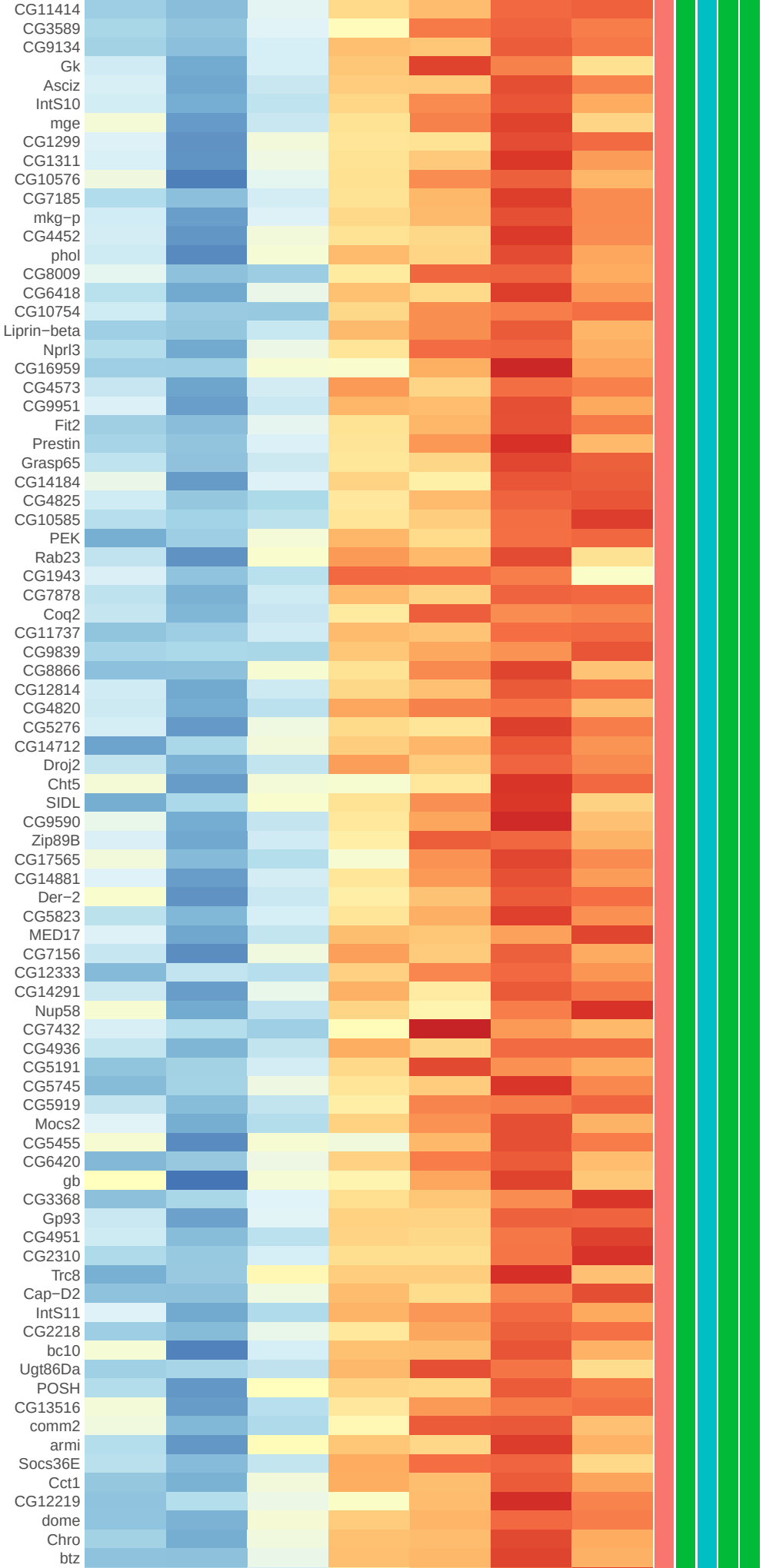
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phol
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CG6418
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Liprin−beta
Nprl3
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Fit2
Prestin
Grasp65
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PEK
Rab23
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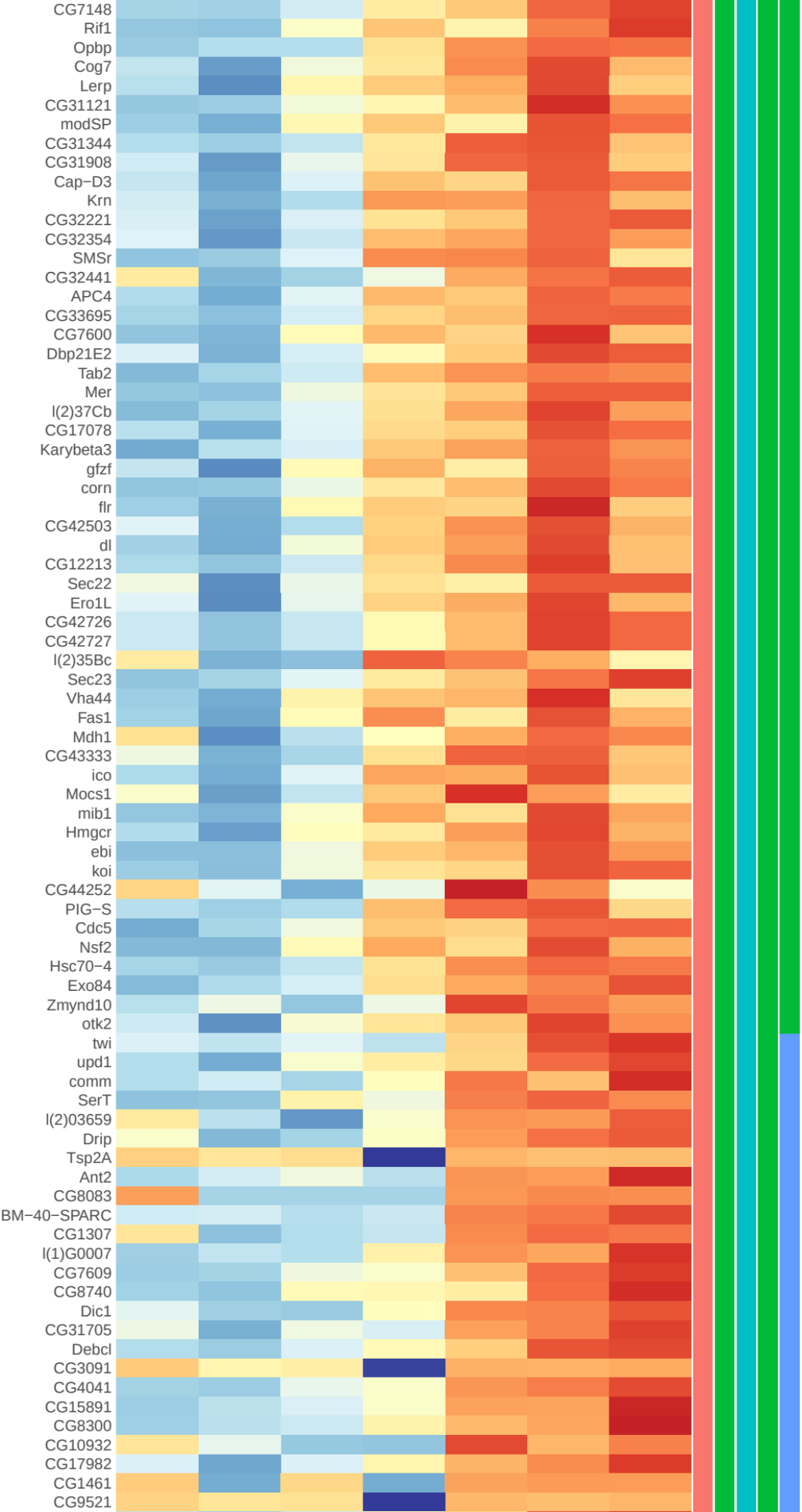
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koi
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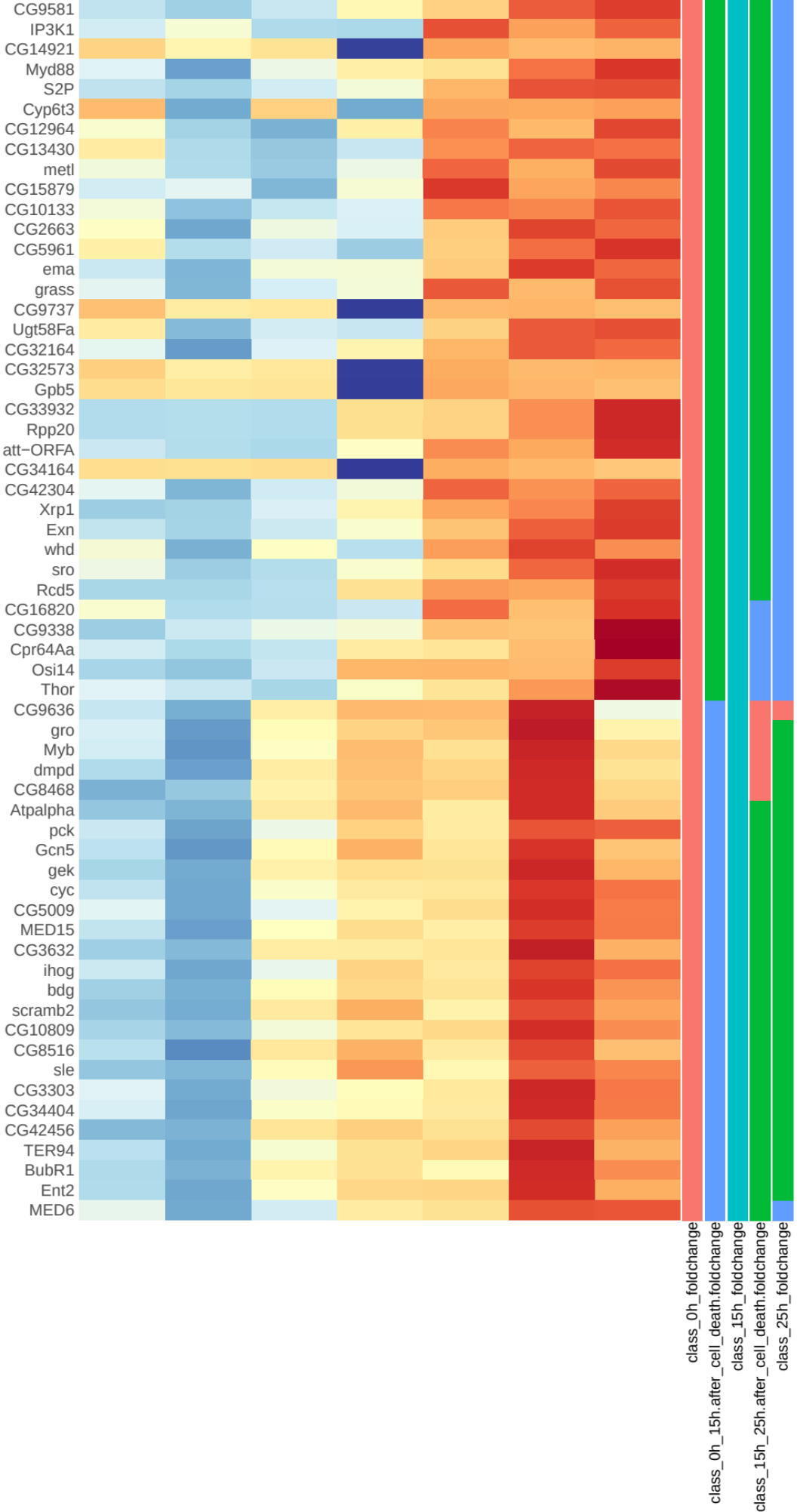
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Exn
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Osi14
Thor
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