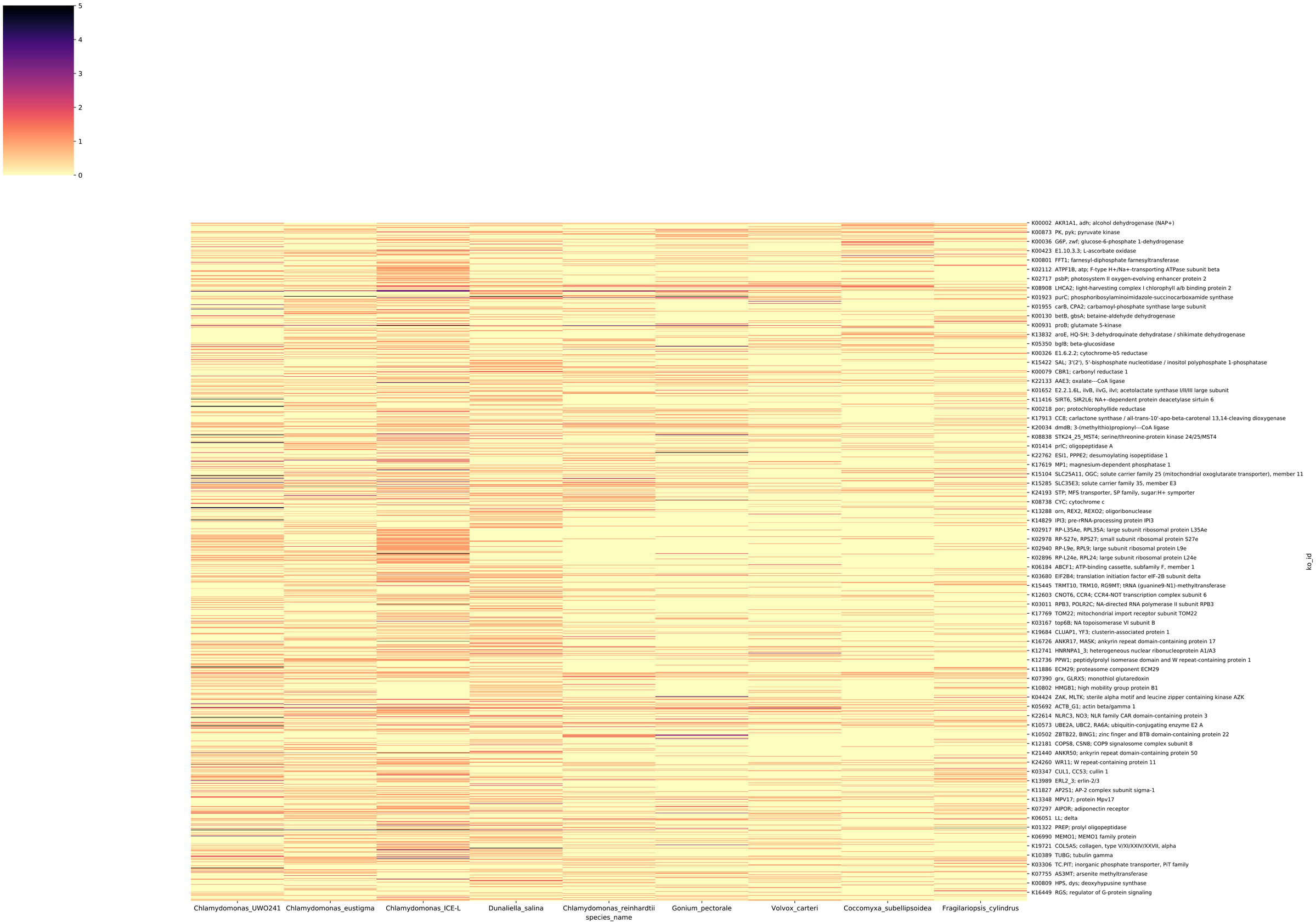
5
4
3
2
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K00002 AKR1A1, adh; alcohol dehydrogenase (NAP+)
K00873 PK, pyk; pyruvate kinase
K00036 G6P, zwf; glucose-6-phosphate 1-dehydrogenase
K00423 E1.10.3.3; L-ascorbate oxidase
K00801 FFT1; farnesyl-diphosphate farnesyltransferase
K02112 ATPF1B, atp; F-type H+/Na+-transporting ATPase subunit beta
K02717 psbP; photosystem II oxygen-evolving enhancer protein 2
K08908 LHCA2; light-harvesting complex I chlorophyll a/b binding protein 2
K01923 purC; phosphoribosylaminoimidazole-succinocarboxamide synthase
K01955 carB, CPA2; carbamoyl-phosphate synthase large subunit
K00130 betB, gbsA; betaine-aldehyde dehydrogenase
K00931 proB; glutamate 5-kinase
K13832 aroE, HQ-5H; 3-dehydroquinate dehydratase / shikimate dehydrogenase
K05350 bglB; beta-glucosidase
K00326 E1.6.2.2; cytochrome-b5 reductase
K15422 SAL; 3'(2'), 5'-bisphosphate nucleotidase / inositol polyphosphate 1-phosphatase
K00079 CBR1; carbonyl reductase 1
K22133 AAE3; oxalate---CoA ligase
K01652 E2.2.1.6L, ilvB, ilvG, ilvI; acetolactate synthase I/II/III large subunit
K11416 SIRT6, SIR2L6; NA+-dependent protein deacetylase sirtuin 6
K00218 por; protochlorophyllide reductase
K17913 CC8; carlactone synthase / all-trans-10'-apo-beta-carotenal 13,14-cleaving dioxygenase
K20034 dmdB; 3-(methylthio)propionyl---CoA ligase
K08838 STK24_25_MST4; serine/threonine-protein kinase 24/25/MST4
K01414 prlC; oligopeptidase A
K22762 ESI1, PPPE2; desumoylating isopeptidase 1
K17619 MP1; magnesium-dependent phosphatase 1
K15104 SLC25A11, OGC; solute carrier family 25 (mitochondrial oxoglutarate transporter), member 11
K15285 SLC35E3; solute carrier family 35, member E3
K24193 STP; MFS transporter, SP family, sugar:H+ symporter
K08738 CYC; cytochrome c
K13288 orn, REX2, REXO2; oligoribonuclease
K14829 IPI3; pre-rRNA-processing protein IPI3
K02917 RP-L35Ae, RPL35A; large subunit ribosomal protein L35Ae
K02978 RP-S27e, RPS27; small subunit ribosomal protein S27e
K02940 RP-L9e, RPL9; large subunit ribosomal protein L9e
K02896 RP-L24e, RPL24; large subunit ribosomal protein L24e
K06184 ABCF1; ATP-binding cassette, subfamily F, member 1
K03680 EIF2B4; translation initiation factor eIF-2B subunit delta
K15445 TRMT10, TRM10, RG9MT; tRNA (guanine9-N1)-methyltransferase
K12603 CNOT6, CCR4; CCR4-NOT transcription complex subunit 6
K03011 RPB3, POLR2C; NA-directed RNA polymerase II subunit RPB3
K17769 TOM22; mitochondrial import receptor subunit TOM22
K03167 top6B; NA topoisomerase VI subunit B
K19684 CLUAP1, YF3; clusterin-associated protein 1
K16726 ANKR17, MA5K; ankyrin repeat domain-containing protein 17
K12741 HNRNPA1_3; heterogeneous nuclear ribonucleoprotein A1/A3
K12736 PPW1; peptidylprolyl isomerase domain and W repeat-containing protein 1
K11886 ECM29; proteasome component ECM29
K07390 grx, GLRX5; monothiol glutaredoxin
K10802 HMGB1; high mobility group protein B1
K04424 ZAK, MLTK; sterile alpha motif and leucine zipper containing kinase AZK
K05692 ACTB_G1; actin beta/gamma 1
K22614 NLRC3, NO3; NLR family CAR domain-containing protein 3
K10573 UBE2A, UBC2, RA6A; ubiquitin-conjugating enzyme E2 A
K10502 ZBTB22, BING1; zinc finger and BTB domain-containing protein 22
K12181 COPS8, CSN8; COP9 signalosome complex subunit 8
K21440 ANKR50; ankyrin repeat domain-containing protein 50
K24260 WR11; W repeat-containing protein 11
K03347 CUL1, CC53; cullin 1
K13989 ERL2_3; erlin-2/3
K11827 AP2S1; AP-2 complex subunit sigma-1
K13348 MPV17; protein Mpv17
K07297 AIPOR; adiponectin receptor
K06051 LL; delta
K01322 PREP; prolyl oligopeptidase
K06990 MEMO1; MEMO1 family protein
K19721 COL5AS; collagen, type V/XI/XXIV/XXVII, alpha
K10389 TUBG; tubulin gamma
K03306 TC.PIT; inorganic phosphate transporter, PiT family
K07755 AS3MT; arsenite methyltransferase
K00809 HPS, dys; deoxyhypusine synthase
K16449 RGS; regulator of G-protein signaling
ko_id
Chlamydomonas_UWO241
Chlamydomonas_eustigma
Chlamydomonas_ICE-L
Dunaliella_salina
Chlamydomonas_reinhardtii
Gonium_pectorale
Volvox_carteri
Coccomyxa_subellipsoidea
Fragilariopsis_cylindrus
species_name