Printolian co-depareme complex, PKS-RHO1 variant[12;1,3-beta-D-glucan synthase complex, PKS-RHO1 variant[12;1,3-beta-D-glucan synthasynthase complex, PKS-RHO1 variant[12;1,3-beta-D-glucan synthase c
Box H/ACA ribonucleoprotein complex/114 1 0 0 1 1 cAMP-dependent protein kinase complex variant 4 13;cAMP-dependent protein kinase complex 4 12;cAMP-dependent protein kinase complex 4 12;cAMP-depende
cAMP-dependent protein kinase complex variant 4 13;cAMP-dependent protein kinase complex variant 5 13;cAMP-dependent protein kinase complex variant 1 12;cAMP-dependent protein kinase complex variant 1 12;cAMP-d
Exomer complex 1 5
MTQ2-TRM112 eRF1 methyltransferase complex/ 12;BUD23-TRM112 methyltransferase complex/ 12;TRM1-TRM112 tRNA (m2G10) methyltransferase complex/ 00 1 0 0 0 0   Nucleotide excision repair factor 4 complex/ 13 1 0 0 0 0   PAN1 actin cytoskeleton-regulatory complex/ 13;SLAC complex/ 12 1 0 0 0 0   Peptide:N-glycanase-Rad23 complex/ 12;CDC48-CDC23-UFD2 complex/ 12;Nucleotide excision repair factor 2 complex/ 12 1 0 0 0 0   PMT1-PMT2 dolichyl-phosphate-mannose-protein mannosyltransferase complex/ 12;PMT1-PMT3 dolichyl-phosphate-mannose-protein mannosyltransferase complex/ 12 1 0 0 0
Nucleotide excision repair factor 4 complex 1 3
PAN1 actin cytoskeleton—regulatory complex 1 3;SLAC complex 1 2 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
Peptide:N-glycanase-Rad23 complex 1 2;CDC48-CDC23-UFD2 complex 1 2;Nucleotide excision repair factor 2 complex 1 2 — 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
Peptide:N-glycanase-Rad23 complex 1 2;CDC48-CDC23-UFD2 complex 1 2;Nucleotide excision repair factor 2 complex 1 2 — 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
PMT1-PMT2 dolichyl-phosphate-mannose-protein mannosyltransferase complex 1 2;PMT1-PMT3 dolichyl-phosphate-mannose-protein mannosyltransferase complex 1 2 0
KAPT-GCK1-GCK2 transcription activation complex[1]5, leiosome[1]5, KAPT-GCK1 transcription activation complex[1]2
$RENTcomplex 1 3 \qquad \qquad 1 \qquad \qquad 0 \qquad \qquad 1$
RNA polymerase I upstream activating factor complex 1 6;Nucleosome, variant HTA1-HTB1 1 4;Nucleosome, varian
RSC complex variant RSC2 1 16;RSC complex variant RSC1 1 16 0 0
SAGA complex 1 20;SLIK (SAGA-like) complex 0  0 0
Ydr131c ubiquitin ligase complex 1 5;SCF-Ufo1 ubiquitin ligase complex
SDS22-GLC7-YPI1 phosphatase complex 1 3;SDS22-GLC7 phosphatase complex 1 2 — 1 0
DNA-directed RNA polymerase II complex 4 17;DNA-directed RNA polymerase I complex 2 10 — 4 1
26S Proteasome complex 13 33 — 11 0
Nuclear pore complex 3 32
Arp2/3 complex 2 7 — 2 0 0
Gin4 serine/threonine kinase complex 2 8;Septin complex 1 5 2 1
INO80 chromatin remodeling complex 2 15;NuA4 histone acetyltransferase complex 1 15;Swr1 chromatin remodelling chrom
Mitochondrial respiratory chain complex III 2 10 2 1
Ribonucleoside-diphosphate reductase variant 2 2 4;Ribonucleoside-diphosphate reductase variant 1 2 3
TIM22 mitochondrial inner membrane twin-pore carrier translocase complex 2 6;TIM9-TIM10 mitochondrial intermembrane space protein transporter complex 2 2 0 1 1 TDEX and the last and the l
TREX complex 2 7 — 2 0 0
U4/U6.U5 tri-snRNP complex 2 28;U4/U6 snRNP 2 17;U6 snRNP complex 2  8;LSM2-8 complex 2  7;LSM2-7 complex 2  6 — 2 1
Mitochondrial pyruvate dehydrogenase complex 3 5;Glycine decarboxylase multienzyme complex 1 4;Mitochondrial 2–oxoglutarate dehydrogenase complex 1 4
COPII vesicle coat complex 3 4;SEC23–LST1 COPII cargo recruitment complex 1 2 0 0
Eukaryotic translation initiation factor 4F complex, variant TIF4631 3 3;Eukaryotic translation initiation factor 4F complex, variant TIF4632 2 3 0 0
GET4-GET5 transmembrane domain recognition complex 3 5 0 0
Eukaryotic translation initiation factor 3 core complex 4 5 4 0 1
Vacuolar proton translocating ATPase complex, vacuole variant 8 14; Vacuolar proton translocating ATPase complex, Golgi variant 7 14 7 0 1 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
40S cytosolic small ribosomal subunit 8 9 — 8 0
HMC complex 6 7;HAP1 transcriptional repressor complex, SSA1 variant 3 4;HAP1 transcriptional repressor complex, SSA2 variant 3 4 — 6 6
60S cytosolic large ribosomal subunit 5 10 — 5 0
Elongation Factor eEF1 complex, variant TEF4 3 3;Elongation Factor eEF1 complex, variant CAM1 3 3 — 4 0
Mitochondrial proton-transporting ATP synthase complex 2 17 2 0 0
Luminal surveillance complex 1 3 — 1 0 0
mRNA cleavage factor complex CF I 1 9 — 1 0 0
Myosin class II complex 1 3;Myosin class V complex, MYO4 variant 1 2;MIH complex 1 2;Myosin class V complex, MYO2 variant 1 1 1 1 1
Ribosome quality control complex 1 6;CDC48-NPL4-UFD1 AAA ATPase complex 1 3;CDC48-NPL4-VMS1 AAA ATPase complex 1 3;CDC48-CDC23-UFD2 complex 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
SET3C histone deacetylase complex 1 7
Spindle pole body central plaque complex MYO5 variant plaque complex, MYO5
Trehalose-6-phosphate synthase/phosphatase variant 2 1 3 1 1 0
Vesicular SNARE complex SSO2–SEC9–SNC2 1 3; Vesicular SNARE complex SSO2–SEC9–SNC1 1 3 1 1 0 0
Chaperonin-containing T-complex 4 8 — 3 1 0
COPI vesicle coat complex 4 7 — 3
TOM40 mitochondrial outer membrane translocase holocomplex 2 7;TOM40 mitochondrial outer membrane translocase core complex 1 5 — 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
Small ribosomal subunit processome, variant 1 8 47;Small ribosomal subunit processome, variant 3 8 47;Box C/D snoRNP complex 3  4;U4/U6.U5 tri–snRNP complex 3  4;U4/U6.snRNP 1  1 — 6 7
TIM23 mitochondrial inner membrane pre-sequence translocase complex, motor variant 3 9;TIM23 mitochondrial inner membrane pre-sequence translocase complex, sort variant 1 5;COX1 pre-assembly complex 1 4;Endonuclease Scel 1 1 2 3
PAF1 complex 2 5 0 0
ESCRT-III complex 1 4 0 0