

Supplementary table S2. List of protein complexes.

The first column provides the complex identifier and the second the name of the complex, if known. The third column contains the core proteins (i.e. those present in 2/3rds of the isoforms). The fourth column includes all the modules associated to this particular complex; numbers refers to the module identifier (see Table S3). The fifth column shows all the attachment proteins found in any of the complex isoforms. Finally, the sixth column reports the most likely subcellular localization of the complex. The complex localization reported corresponds to the most common of those for each individual core components as defined in Huh *et al.*³⁶. Localizations. Bd: Bud. Cy: Cytoplasmic. ER: Endoplasmic Reticulum. Go: Golgi. Mt: Mitochondrial. No: Nucleolar. Nu: Nuclear. Va: Vacuolar. NP: Nuclear Periphery. CP: Cell Periphery. Cs: Cytoskeleton.

ID	Complex Name	Core*	Modules**	Attachments	Localizations
1	U1 snRNP	Luc7 Mud1 Nam8 Prp39 Prp40 Prp42 Snp1 Snu56 Snu71 Yhc1	7: <i>Mud1 Nam8 Prp40 Prp42 Snp1 Snu56 Snu71 Sto1 Yhc1</i> 13: Prp3 Prp31 Smb1 Spp381 74: Cbc2 Npl3 83: <i>Prp39 Smd1</i> 106: <i>Luc7 Smd3</i> 122: Air2 Nab3 Nrd1 Trf4 123: Bur2 Sgv1 142: Cus1 Hsh155 Lea1 Prp21 Rse1	Aar2 Brr1 Brr2 Csn12 Dib1 Hsh49 Hta1 Msl1 Mtr4 Pab1 Pan2 Pho80 Prp11 Prp38 Prp4 Prp6 Prp8 Prp9 Rpl16b Rps22a Scp160 Sen1 Smd2 Snt309 Snu114 Snu23 Snu66 Sro9 Tif4631 Tif4632 Ybl104c Ylr424w	Nu
2	20S core particle of the proteasome	Pre10 Pre2 Pre3 Pre6 Pre8 Pup3 Scf1	57: <i>Pre4 Pre8</i> 93: Nas6 Rpn3 141: <i>Pre10 Pre2 Pre3 Pre6 Pup3 Scf1</i>	Blm3 Ecm29 Imp4 Mlh2 Pre1 Pre5 Pre7 Pre9 Rpn10 Rpn2 Rpn8	Cy, Nu
3	Lsm2-Lsm8 complex	Lsm2 Lsm4 Lsm6 Lsm7 Pat1	1: <i>Lsm6 Lsm7 Prp4 Prp6 Prp8 Snu114</i> 13: Prp3 Prp31 Smb1 Spp381 145: Lsm1 <i>Lsm2 Lsm4 Lsm5 Pat1</i>	Aar2 Brr1 Brr2 Bud31 Cef1 Cwc2 Cwc22 Cwc23 Dhh1 Dib1 Gcn2 Kem1 Krs1 Lin1 Mrp18 Msl1 Nam7 Pab1 Prp24 Prp38 Prp9 Smd1 Smd2 Smd3 Snu23 Snu66 Ybl104c Ykr022c Ylr424w	Cy, Nu
4	Casein kinase II complex	Cka1 Cka2 Ckb1 Ckb2	18: Ppz1 Ykl088w	Cdc73 Chd1 Ctr9 Hhf1 Hot1 Leo1 Nip1 Paf1 Pob3 Prt1 Rok1 Rpg1 Rrp7 Rtf1 Spt16 Ssa3 Ssa4 Utp22 Ylr003c	Cy, Nu
5	Complex 5	Dbp2 Rpl2a	4: Lsg1 Nmd3 Rei1	Arx1 Cdc123 Mis1 Pim1 Rpl11b Rpl12b Rpl13a Rpl16b Rpl17a Rpl18a Rpl20b Rpl27a Rpl27b Rpl33a Rpl36a Rpl36b Rpl6b Rpl8b Rpp0 Rps14b Shm1 Ypl009c	Cy
6	Complex 6	Ecm5 Psd2 Yfr006w Yol054w	96: Hhf1 <i>Yol054w</i>	Cdc55 Iml1 Kre33 Mdn1 Mis1 Nmd5 Pro1 Rpl10 Rpl13b Rpl15a Rpl38 Rps11b Rps9a Rps9b Uso1 Xks1	Cy, Nu
7	Complex 7	Nsa1 Rpf1	17: Has1 <i>Nsa1 Rpf1</i> 49: Brx1 Erb1 Nop15 Nop7 52: Cic1 Ipi1 61: Noc2 Nop12 Rrp5 88: Dbp9 Nop4 95: Loc1 Rpl9a	Drs1 Ebp2 Mak21 Mak5 Mrt4 Noc3 Nog1 Nsa2 Rix1 Rix7 Rlp7 Rpl12a Rpl13a Rpl15a Rpl16a Rpl18b Rpl20b Rpl23b Rpl26b Rpl30 Rpl31a Rpl4b Rpl5 Rpl7a Rpl8b Rpp2b Rrp1 Rsa3 Spb1 Spb4 Tif6 Tsl1 Yor1 Ytm1	Cy, No, Nu
8	m-AAA protease complex	Afg3 Phb2 Yta12	125: <i>Afg3 Phb1 Phb2 Yta12</i>	Gdh1 Hmg1 Kap122 Pgi1 Pmr1	Mt
9	Mitochondrial ribosomal small subunit	Ehd3 Mrp1 Mrp13 Mrp17 Mrp21 Mrp4 Mrp51 Mrps16 Mrps17 Mrps18 Mrps28 Mrps5 Mrps8 Mrps9 Nam9 Pet123 Rsm10 Rsm19 Rsm22 Rsm23 Rsm25 Rsm27 Rsm7	59: <i>Mrp1 Rsm25</i> 73: Mrps35 <i>Nam9</i> 112: Pus7 Tan1	Ade6 Ded1 Met6 Psd2 Rsm24 Rsm26 Ubp10	Mt
10	Complex 10	Mae1	60: Acs2 Rnr1 63: <i>Mae1</i> Tom1 81: Gcn1 Gcn20 100: Rvb2 Sec53	Acc1 Ade3 Aro1 Aro3 Cct4 Cct6 Cse1 Dps1 Eft1 Eft2 Fas1 Fas2 Fur1 Gea2 Gfa1 Grs1 Hsc82 Hsp60 Hsp82 Iis1 Ilv1 Ilv5 Mdn1 Msn5 Pfk1 Prs3 Psa1 Rpl18b Rpl30 Rps0a Sap155 Sar1 Sec21 Tdh1 Tfp1 Trm3 Tub1 Tub3 Ura7 Vps13 Ydr341c Yer036c Yhb1 Ynk1	Cy, Mt, Nu
11	Complex 11	Nob1	79: <i>Nob1</i> Rps18b 102: Rps9b Slx9	Bem2 Emg1 Enp1 Imp3 Imp4 Krr1 Mak5 Noc4 Pno1 Pwp2 Rio2 Rpl30 Rpl31a Rps10a Rps11b Rps13 Rps14a Rps17b Rps21a Rps22b Rps24a Rps24b Rps28b Rps8a Snf2 Sof1 Tsr1 Utp6 Ygr054w	Cy, No, Nu
12	Complex 12	Mot1	37: Ioc3 Isw1 <i>Mot1</i>	Abf2 Hhf1 Hmo1 Hta2 Net1 Pdr1 Reb1 Ret1 Rpl20a Rps3 Rsc3 Rvb1 Sec2 Spt15 Taf1 Tao3 Top1 Vps1 Yil056w	Nu
13	Trehalose-6-phosphate synthase/phosphatase complex	Gln1 Mcm2 Mks1 Tsc10	50: Tps1 Tps2 Tps3	Hsp42 Iml1 Reg1 Rpl10 Tif1 Tif2 Tsl1	Cy
14	V0 vacuolar ATPase complex	Emi2 Vma6 Vph2	41: <i>Vma6 Vph2</i>	Bni1 Drs2 Gaa1 Lys9 Nop6 Pdc1 Pgi1 Vac8 Vma10 Vma2 Vma7 Vph1 Vtc4 Yor1 Yra1	Cy, Va
15	Complex 15	Rps25a Rrn3		Erg13 Rpl13b Rpl17a Rpl24a Rpl25 Rpl30 Rpl35a Rpl7a Rps22a Rps24b Tef2	Cy, No, Nu
16	Complex 16	Pph3 Psy2 Ybl046w	53: Spt4 Spt5	Bem2 Rga2 Tdh3 Trm3	Cy, Nu
17	Complex 17	Bcp1 Ypl208w		Eft1 Gfa1 Glt1 Kap123 Rpl23a Rpl23b Rpl6a Rps5 Ssa4 Ura2 Yol098c	Cy, Nu

18	Complex 18	Hca4	65: Bms1 Rcl1 72: Hca4 Nop58 Sik1 78: Kre33 Ygr145w 80: Bud21 Imp3	Cbf5 Dbp8 Dim1 Emg1 Ero1 Esf1 Fun12 Gbp2 Imd2 Krr1 Mis1 Mpp10 Mvd1 Nip1 Nop1 Nop12 Nop4 Nop6 Pre9 Prp43 Pwp2 Ref2 Rpl11b Rpl13b Rpl16b Rpl26b Rpl2b Rpl36a Rpl6b Rpp2b Rps16a Rps22a Rps28b Rps4a Rps4b Rps7a Rps9b Rrp5 Rrp9 Sof1 Sro9 Utp10 Utp15 Utp20 Utp30 Utp7 Utp9 Ylr003c Ynl224c	Cy, No
19	Complex 19	Ist2		Ded81 Glt1 Hir2 Mrpl7 Rpl10 Rpl11a Rpl13b Rpl17b Rpl18a Rpl24a Rpl25 Rpl26b Rpl28 Rpl2b Rpl33a Rpl38 Rpl7a Rps14a Rps16b Rps18a Rps20 Rps4a Rps5 Rps6b Rps9a Tif34	Cy
20	Complex 20	Loc1 Rpl9a	95: Loc1 Rpl9a	Arx1 Cin1 Dbp10 Drs1 Erb1 Fpr4 Lhp1 Mak5 Mrt4 Nip7 Noc2 Noc3 Nog1 Nop15 Nop2 Nop7 Nsa2 Nug1 Rlp7 Rpf2 Rpl11b Rpl13a Rpl13b Rpl16a Rpl2b Rpl3 Rpl43a Rrp1 Sda1 Spb4 Tif6 Ytm1	Cy, No, Nu
21	Complex 21	Shm1	113: Fun12 Rpl12b	Ade3 Bni1 Cbf5 Dbp2 Mis1 Puf6 Rat1 Rpl10 Rpl13a Rpl16b Rpl17a Rpl18a Rpl24a Rpl25 Rpl27a Rpl27b Rpl28 Rpl2a Rpl2b Rpl33a Rpl6a Rps14b Rps1a Rps22b Ypl009c	Cy, Mt, Nu
22	Sin3 Histone deacetylase complex	Cti6 Rpd3 Rxt3 Sap30 Sds3 Sin3 Ume1 Ymr075w	129: Arp4 Eaf3 Epl1 Esa1 Vid21	Abf2 Cka1 Cka2 Ckb1 Dep1 Eaf5 Ecm16 Hta2 Ioc3 Isw1 Pho23 Rpl17a Rpl33a Rpl35b Rsc8 Spt6 Top1 Tra1 Tup1 Ume6 Yaf9 Ybr095c	Nu
23	Complex 23	Emg1 Kre33 Krr1 Noc4 Nop14 Ygr145w	38: Emg1 Noc4 Nop14 Utp22 62: Imp4 Rrp9 Utp13 65: Bms1 Rcl1 69: Ecm16 Krr1 Nop6 78: Kre33 Ygr145w 79: Nob1 Rps18b 80: Bud21 Imp3 102: Rps9b Slx9	Bfr2 Cbf5 Dbp8 Enp1 Ero1 Esf1 Fap7 Kri1 Mpp10 Mvd1 Nop1 Pre9 Rok1 Rpl15a Rpl19a Rps10b Rps11b Rps13 Rps14a Rps16a Rps17b Rps1b Rps20 Rps22a Rps22b Rps24b Rps7a Rps8a Rrp12 Rrp5 Rrp7 Sof1 Tsr1 Utp20 Utp30 Utp6 Ylr003c	Cy, No, Nu
24	Complex 24	Bud20	47: Nog2 Nsa2 Nug1 Tif6 133: Arx1 Bud20 Ipi3 Rix1 Sda1 Ycr072c	Mrt4 Nog1 Nop15 Nop7 Rlp7 Rpf2 Rpl11a Rpl11b Rpl25 Rpl27a Rpl27b Rpl28 Rpl2b Rpl3 Rpl30 Rpl31a Rpl31b Rpl36b Rpl38 Rpl7a Rpl8a	Cy, Nu
25	DNA repair complex	Mph1 Msh6 Rad52 Rfa1	67: Msh2 Msh3 70: Rad16 Rim1	Dna2 Kap95 Lcd1 Mec1 Pkg1 Rad51 Rfa2 Rfa3 Vps1 Ydl156w	Cy, Nu
26	Complex 26	Met18 Rad3		Atp2 Cdc19 Pkg1 Rpt6 Tfp1 Tub1 Ydj1	Cy, Nu
27	Complex 27	Nop14 Utp8	16: Enp1 Hrr25 Ltv1 Rio2 Tsr1 26: Nan1 Utp10 Utp15 Utp4 Utp7 Utp9 28: Pwp2 Utp18 Utp21 Utp6 38: Emg1 Noc4 Nop14 Utp22 62: Imp4 Rrp9 Utp13 65: Bms1 Rcl1 69: Ecm16 Krr1 Nop6 72: Hca4 Nop58 Sik1 78: Kre33 Ygr145w 80: Bud21 Imp3 86: Dim1 Rrp12 102: Rps9b Slx9	Bfr2 Cbf5 Dbp8 Dip2 Ero1 Esf1 Fap7 Gbp2 Hsl1 Kri1 Mpp10 Mrd1 Mvd1 Nob1 Nop1 Pno1 Rok1 Rpl15a Rpl16b Rpl17a Rpl19a Rpl26b Rpl33a Rpl36a Rpl6b Rpp2b Rps11b Rps13 Rps14a Rps14b Rps16a Rps17a Rps17b Rps18a Rps1b Rps21a Rps22b Rps24b Rps25b Rps28b Rps3 Rps4a Rps4b Rps7a Rps8a Rps8b Rps9a Rrp7 Sof1 Sro9 Utp20 Utp30 Ygr054w Ylr003c Yor059c	Cy, No, Nu
28	Complex 28		5: Img1 Mhr1 Mrp49 Mrpl9 Yml6 45: Mrpl10 Mrpl16 Mrpl28 Mrpl7 118: Mrpl39 Mrpl44 127: Ede1 Syp1	Aep1 Atg26 Bud3 Cop1 Img2 Mrp20 Mrp7 Mrpl1 Mrpl13 Mrpl15 Mrpl17 Mrpl20 Mrpl22 Mrpl23 Mrpl24 Mrpl27 Mrpl3 Mrpl35 Mrpl36 Mrpl4 Mrpl6 Mrpl8 Pet56 Ret2 Rom2 Sec26 Sec28 Shq1 Sro9 Utp20 Ymr098c Ypl183w-a	Go, Mt
29	Gcn1/Gcn2/Gcn20 complex	Gcn1 Gcn20	81: Gcn1 Gcn20 92: Cse1 Sec7 100: Rvb2 Sec53	Aha1 Arf1 Cct8 Fur1 Gfa1 Idh2 Lys12 Msn5 Pet9 Sst2 Tub1 Ufd4 Vps13	Cy, Nu
30	RNA polymerase I	Rpa12 Rpa190 Rpa43 Rpa49 Rpc40	121: Rpa12 Rpa135 Rpa190 Rpa43 Rpb5 Rpc40	Ams1 Cdc14 Dhh1 Hsp42 Met10 Rpb10 Rpb8 Rpe31 Rpo26 Sro7 Yer139c	No, Nu
31	mRNA decapping complex	Dcp1 Dcp2 Edc3	128: Dcp1 Dcp2 Edc3	Kem1 Rpm2 Ybr094w	Va
32	Ribosome-associated complex (RAC)	Scp160 Ssz1 Zuo1	105: Ski2 Ski3 Ski8 109: Rpl8a Yer036c 113: Fun12 Rpl12b 120: Rai1 Rat1	Ade5,7 Amn1 Asc1 Bem2 Cdc25 Mis1 Rpl10 Rpl11a Rpl25 Rpl27a Rpl3 Rpl30 Rpl6a Rpl8b Rps10a Rps11a Rps13 Rps14b Rps17b Rps20 Rps21a Rps22a Rps3 Rps4a Rps5 Rps7a Tif4631 Yal036c Ygr054w	Cy

33	U2 snRNP	Cus1 Hsh155 Hsh49 Lea1 Msl1 Prp11 Prp21 Prp9 Rse1	9: Cwc23 Prp43 Ylr424w 13: Prp3 Prp31 Smb1 Spp381 106: Luc7 Smd3 142: <i>Cus1 Hsh155 Lea1 Prp21 Rse1</i>	Brr1 Brr2 Cdc33 Clf1 Csn12 Dib1 Isy1 Lin1 Prp38 Smd1 Smd2 Snt309 Snu23 Snu66 Syf1 Ybl104c Yhc1	Nu
34	Scs2/Opi1 complex	Swh1	40: Mrp8 Pil1	Num1 Opi1 Osh2 Rps4b Rps7a Scs2 Ser33 Sup45 Yar044w Ygr130c	Cy, Va
35	Sit4/Sap190 complex	Sap190	51: Cdc25 Mds3	Acc1 Cct2 Cct5 Dig1 Fab1 Fyy8 Mrp13 Rpl11b Sap155 Sit4	Cy, Nu, Va
36	Complex 36	Fur1 Sar1 Ydr341c Ynk1	116: Hsp82 Lys12 139: Cop1 Ret2 Ret3 Sec26 Sec27 Sec28	Aap1' Ade3 Aro1 Cdc19 Cse1 Dps1 Eft2 Fas2 Gcn20 Grs1 Hsc82 Ilv5 Ilv6 Msn5 Pfk1 Rpa135 Rpl30 Rps0a Rpt5 Sec21 Sec53 Tdh1 Tfp1 Tom1 Trm3 Vps13	Cy, Nu
37	Complex 37		28: Pwp2 Utp18 Utp21 Utp6 38: Emg1 Noc4 Nop14 Utp22 62: Imp4 Rrp9 Utp13 78: Kre33 Ygr145w 80: Bud21 Imp3 102: Rps9b Slx9	Bms1 Cbf5 Ckb1 Dim1 Dip2 Ecm16 Enp1 Ero1 Esf1 Hsl1 Krr1 Ltv1 Mpp10 Mvd1 Nan1 Nop58 Pno1 Rok1 Rpl17a Rpl1b Rpl27a Rpl30 Rpl36a Rps13 Rps16a Rps17b Rps18a Rps22b Rps24a Rps25b Rps3 Rps5 Rps7a Rps8a Rps9a Rtg2 Sec21 Sik1 Sof1 Tif1 Tsr1 Utp10 Utp15 Utp20 Utp30 Utp7 Utp9 Ylr003c Ynk1	Cy, No, Nu
38	RNA polymerase III complex	Ret1 Rpc25 Rpc31 Rpc34 Rpc37 Rpc40 Rpc53 Rpc82 Rpo31	121: Rpa12 Rpa135 Rpa190 Rpa43 Rpb5 <i>Rpc40</i>	Ams1 Arp8 Cdc14 Dhhl1 End3 Fba1 Hta2 Htb1 Maf1 Reb1 Rpa49 Rpb10 Rpb8 Rpe11 Rpe17 Rpe19 Rpo26 Rsc30 Rsc8 Sro7 Tfc4 Yil056w	Nu
39	Complex 39	Bud20 Ipi3 Nog1 Nog2 Nsa2 Nug1 Rix1 Rpf2 Sda1 Ycr072c	4: Lsg1 Nmd3 Rei1 47: <i>Nog2 Nsa2 Nug1 Tif6</i> 49: Brx1 Erb1 Nop15 Nop7 52: Cic1 Ipi1 91: Ebp2 <i>Rpf2</i> 95: Loc1 Rpl9a 133: <i>Arx1 Bud20 Ipi3 Rix1 Sda1 Ycr072c</i>	Cin1 Dbp10 Drs1 Mdn1 Mrt4 Nip7 Noc2 Noc3 Rlp7 Rpl11a Rpl12a Rpl16a Rpl18a Rpl20a Rpl20b Rpl25 Rpl27a Rpl28 Rpl3 Rpl30 Rpl31a Rpl31b Rpl33a Rpl36a Rpl36b Rpl38 Rpl43a Rpl5 Rpl7a Rrp1 Spb1 Spb4	No, Nu
40	Cyr1/Srv2 complex	Cyr1 Pho81 Srv2		Act1 Mrp151 Pcl8 Pho80 Pho85 Ygr250c	Cy, Nu
41	Complex 41	Dbp8 Esf1 Mvd1		Bfr2 Bud21 Cbf5 Dip2 Emg1 Ero1 Hca4 Krr1 Mpp10 Nop58 Pre9 Rok1 Rpl11b Rpl15a Rpl16b Rpl19a Rpl1b Rpl26b Rpl2b Rpl33a Rpl36a Rpl4a Rpl6b Rpp0 Rpp2b Rps14a Rps14b Rps16a Rps17b Rps18a Rps1b Rps22a Rps24b	Cy, No, Nu
42	Cytoskeleton-regulatory Lsb3/Las17 complex	Las17 Lsb3 Sla1	127: Ede1 Syp1 138: Chc1 Clc1	Apm2 Bzz1 Ecm25 End3 Rpl11a Rpl12a Rpl19b Rpl28 Rpl35a Rpl38 Rpl6b Rpl9a Rps11a Rps14b Sla2 Stm1 Ydr348c Ypt7	Cs, Cy, Go, Va
43	Nup82/Nsp1/Nup159 nuclear pore subcomplex	Nsp1 Nup159 Nup82		Gle2 Mrp11 Mrp115 Mrp124 Mrp13 Mrp135 Mrp18 Nic96 Nup116 Nup192 Nup57 Ssa2	NP
44	Actin-associated motorproteins 1 complex	Myo2 She4	33: <i>Myo2 She4</i> 58: Mlc1 Myo1 68: Cmd1 Myo5	Arr4 Hsc82 Mam33 Nmd5 Pdr5 Yfl034w	Bd, CP, Cy
45	Translation initiation factor eIF3 complex	Fun12 Hcr1 Nip1 Prt1 Rli1 Rpg1 Tif34 Tif35 Tif5		Atg9 Cbf5 Ded1 Gcd11 Glt1 Ist2 Rpl13a Rpl16a Rpl16b Rpl18a Rpl18b Rpl6a Rpl7a Rpl8b Rps10a Rps10b Rps11a Rps13 Rps14a Rps16b Rps17a Rps18a Rps1b Rps22a Rps22b Rps28b Rps5 Rps6a Rps6b Rps7a Rps7b Rps8a Rps8b Rps9a Sui1	Cy
46	Clathrin	Chc1 Clc1	101: Sse1 Tsa1 138: <i>Chc1 Clc1</i>	End3 Fas1 Las17 Sla2 Yhb1	Cy, Go
47	Complex 47	Nap1 Nis1 Yol070c	77: Mss51 Ssc1 117: Gin4 Kcc4	Hmo1 Kap114 Pgi1 Rps1a Trm3 Yra1	Bd, CP, Cy
48	Complex 48	Nop8 Urb1 Urb2	124: <i>Urb1 Urb2</i>	Arg1 Leu4 Leu9 Mak5 Nop2 Pre1 Rpl11b Rpl25 Rpl36b Rsa3	No
49	Actin-associated motorproteins 2 complex	Mlc1 Myo4 She2 She3	33: Myo2 She4 58: <i>Mlc1 Myo1</i> 68: Cmd1 Myo5 90: Cdc47 Erg26 131: <i>Myo4 She2 She3</i> 135: Cap1 Cap2 Yir003w	Act1 Brx1 Cdc46 Crn1 Eht1 Hir3 Imd4 Mam33 Nop12 Pdr5 Pol5 Pwp1 Rad53 Yer071c	Bd, CP, Cy, No, Nu
50	Complex 50	Lsg1 Nmd3 Rei1	4: <i>Lsg1 Nmd3 Rei1</i>	Arx1 Bud20 Ipi1 Ipi3 Mdn1 Mis1 Mrt4 Nog1 Nog2 Pim1 Rlp7 Rpl11a Rpl12a Rpl17a Rpl20a Rpl20b Rpl24a Rpl27a Rpl3 Rpl31a Rpl36a Rpl36b Rpl5 Rpl6b Rpl8b Sda1 Tif6 Ycr072c	Cy, Nu

51	Complex 51	Gdb1 Mir1	22: Lcb1 Lcb2 94: Kap104 <i>Mir1</i> 110: Hsp60 Mkt1	Ade5,7 Crm1 Ecm29 Fas1 Gcn1 Gea2 Hsc82 Mpm1 Mtr10 Nmd5 Nup192 Pdr5 Ppz1 Sam2 Tsc13 Tub1 Usa1 Ydj1 Ygr130c	Cy, Nu
52	Signal recognition complex	Sec65 Srp14 Srp54 Srp68 Srp72		Aha1 Bem2 Hsp82 Lhp1 Srp21 Sst2	ER
53	Erv41/Erv46 complex	Ero1 Erv41 Erv46	42: <i>Erv41 Erv46</i>	Fks1 Rot2	
54	Complex 54		9: Cwc23 Prp43 Ylr424w 83: Prp39 Smd1 142: Cus1 Hsh155 Lea1 Prp21 Rse1	Aar2 Brr1 Brr2 Bud31 Cdc40 Csn12 Cwc2 Dib1 Ecm2 Isy1 Lin1 Lsm4 Msl1 Prp11 Prp19 Prp4 Prp45 Prp46 Prp9 Seh1 Smb1 Smd3 Snt309 Snul14 Snu23 Snu66 Syf1 Ybl104c	Cy, Nu
55	Complex 55	Loc1	17: Has1 Nsa1 Rpf1 43: Mak21 Mak5 Nop2 Ytm1 47: Nog2 Nsa2 Nug1 Tif6 48: Fpr4 Puf6 Spb4 49: Brx1 Erb1 Nop15 Nop7 52: Cic1 Ipi1 61: Noc2 Nop12 Rrp5 87: Nip7 Rpl23a 88: Dbp9 Nop4 91: Ebp2 Rpf2 95: <i>Loc1</i> Rpl9a	Arx1 Dbp10 Drs1 Ist2 Lhp1 Mrt4 Noc3 Nog1 Nop1 Rix1 Rix7 Rlp7 Rpl11a Rpl11b Rpl12a Rpl13a Rpl13b Rpl14a Rpl16a Rpl18a Rpl19a Rpl23b Rpl27a Rpl2b Rpl3 Rpl31a Rpl31b Rpl33a Rpl36b Rpl4b Rpl5 Rrp1 Rrp12 Sda1 Spb1 Yer139c	Cy, No, Nu
56	Complex 56	Brx1 Erb1 Mak21 Noc2 Nop4 Puf6	4: Lsg1 Nmd3 Rei1 17: Has1 Nsa1 Rpf1 43: <i>Mak21</i> Mak5 Nop2 Ytm1 47: Nog2 Nsa2 Nug1 Tif6 48: Fpr4 <i>Puf6</i> Spb4 49: <i>Brx1 Erb1</i> Nop15 Nop7 52: Cic1 Ipi1 61: <i>Noc2</i> Nop12 Rrp5 87: Nip7 Rpl23a 88: Dbp9 <i>Nop4</i> 91: Ebp2 Rpf2 95: Loc1 Rpl9a 133: Arx1 Bud20 Ipi3 Rix1 Sda1 Yer072c	Dbp10 Drs1 Mdn1 Mrt4 Noc3 Nog1 Nop1 Nop13 Pol5 Rix7 Rlp7 Rpl11a Rpl11b Rpl12a Rpl13a Rpl13b Rpl14a Rpl15a Rpl15b Rpl16a Rpl19a Rpl1a Rpl1b Rpl20a Rpl23b Rpl25 Rpl26b Rpl27a Rpl27b Rpl28 Rpl2b Rpl3 Rpl31a Rpl31b Rpl33a Rpl33b Rpl35b Rpl36b Rpl38 Rpl43a Rpl5 Rpl6a Rpl6b Rpl7a Rpl7b Rpl8b Rpp2b Rps14a Rps16b Rps18b Rrp1 Rrp12 Rsa3 Spb1 Utp20 Yor1	No, Nu
57	NuA4 Histone Acetyltransferase Complex	Epl1 Esa1	85: Med6 Yap1 129: Arp4 Eaf3 <i>Epl1 Esa1</i> Vid21	Act1 Dep1 Eaf5 Hfi1 Med1 Nam7 Nut1 Pho23 Rpd3 Sgf29 Sgf73 Sin3 Sin4 Snf12 Spt20 Spt3 Spt7 Spt8 Srb2 Taf12 Tra1 Ume1 Yaf9 Ybr095c Ymr075w	Cy, Nu
58	Complex 58	Hrr25 Ltv1 Nob1 Rio2 Tsr1	16: Enp1 <i>Hrr25 Ltv1 Rio2 Tsr1</i> 65: Bms1 Rcl1 79: <i>Nob1</i> Rps18b 86: Dim1 Rrp12 102: Rps9b Slx9 107: Asc1 Ynl224c	Bem2 Dbp2 Dip2 Ecm16 Gen2 Iki3 Kri1 Mpp10 Mrd1 Nop14 Pno1 Rpm2 Rps0a Rps11a Rps11b Rps13 Rps17a Rps17b Rps21a Rps22a Rps22b Rps24b Rps28b Rps3 Rps4a Rps4b Rps7b Rps8a Rps8b Rps9a Snf2 Utp20 Utp22 Ygr054w	Cy, Nu
59	V1 vacuolar ATPase complex	Vma13 Vma4 Vma5 Vma7 Vma8	31: Rav1 Rav2 41: Vma6 Vph2 75: Vma2 <i>Vma5</i>	Emi2 Nop6 Tfp1 Vma10 Vph1 Yer036c	Va
60	Mitochondrial ribosomal large subunit	Atg26 Bud3 Mrpl10 Mrpl16 Mrpl23 Mrpl4 Mrpl51	5: Img1 Mhr1 Mrp49 Mrpl9 Yml6 45: <i>Mrpl10 Mrpl16</i> Mrpl28 Mrpl7 118: Mrpl39 Mrpl44	Aep1 Ded1 Img2 Iml1 Mrp20 Mrp7 Mrpl1 Mrpl13 Mrpl15 Mrpl17 Mrpl20 Mrpl22 Mrpl24 Mrpl27 Mrpl3 Mrpl35 Mrpl36 Mrpl6 Mrpl8 Pet56 Sec1 Shq1 Utp20 Ymr098c Ypl183w-a	Mt
61	Arginine-specific carbamoyl-phosphate synthase complex	Cpa2	55: Cpa1 <i>Cpa2</i>	Aro1 Cct8 Fas1 Gfal1 Gus1 Ilv1 Pro1 Psa1 Rad51 Rpa135 Rpl18b Rps10a Rpt3 Rvb2 Yhb1	Cy
62	Complex 62	Cic1 Ebp2 Ipi1 Rpl31b	52: <i>Cic1 Ipi1</i> 76: Rpo21 Yer139c 91: <i>Ebp2</i> Rpf2	Clu1 Drs1 Mdn1 Mrt4 Nog1 Nog2 Nop15 Rix7 Rpl12a Rpl14a Rpl23b Rpl27a Rpl28 Rpl3 Rpl33a Rrp1 Rrp12 Spb1	No, Nu
63	U4/U6.U5 tri-snRNP	Prp3 Prp31 Prp4 Prp6 Snu66 Spp381	1: Lsm6 Lsm7 <i>Prp4 Prp6</i> Prp8 Snu114 13: <i>Prp3 Prp31</i> Smb1 <i>Spp381</i> 83: Prp39 Smd1 142: Cus1 Hsh155 Lea1 Prp21 Rse1	Aar2 Brr2 Cwc23 Dib1 Lin1 Lsm2 Lsm4 Prp24 Prp38 Prp9 Smd2 Smd3 Snu23 Snu56 Snu71 Yhc1 Ykr022c Ylr424w	Cy, Nu
64	Complex 64	Mtr4 Nab3 Nrd1 Sen1 Sto1	35: Cdc33 Tif4632 74: Cbc2 Npl3 122: Air2 <i>Nab3 Nrd1</i> Trf4 123: Bur2 Sgv1	Brr1 Hta1 Iml1 Lhp1 Nab6 Nam7 Nrp1 Pab1 Rnt1 Rpl4b Sbp1 Sep160 Sgn1 Smd2 Smd3 Sro9 Tif4631	Nu
65	DNA repair complex	Rad16	67: Msh2 Msh3 70: <i>Rad16</i> Rim1	Lcd1 Pep4 Rad52 Vps1 Wrs1 Ydl156w Ymr315w	Cy, Nu, Va

66	Complex 66	Mae1 Prs3 Tom1	63: Mae1 Tom1 81: Gcn1 Gcn20	Acc1 Aha1 Aro1 Cct4 Cct6 Clu1 Crm1 Ded81 Ecm29 Eft1 Fur1 Gea2 Gfa1 Hsc82 Ils1 Ilv1 Kap122 Mdn1 Msn5 Mtr10 Pre6 Pse1 Rpb2 Rpn10 Rpn2 Rpn9 Rpt4 Sam1 Ths1 Tub1 Tub3 Ufd4 Ura7	Cy, Nu
67	Pab1/eIF4G/eIF4E complex	Cdc33 Sbp1 Tif4631 Tif4632	35: Cdc33 Tif4632 98: Cbf5 Ymr310c 115: Sgn1 Ygr250c	Aar2 Brr1 Hca4 Imd3 Imd4 Kem1 Lhp1 Luc7 Mrp7 Nab6 Nam7 Npl3 Nrd1 Nrp1 Pab1 Pol2 Prp43 Pus7 Rpl31b Rps10b Rps16b Rps20 Rps22a Rps22b Sen1 Snp1 Sro9 Sto1 Sxm1 Tif2 Yal036c Ynl224c	Cy, Nu
68	ISW1/IOC3 complex	Ioc3 Isw1	3: Dpb2 Pol2 32: Tbf1 Vid22 37: Ioc3 Isw1 Mot1 66: Hta2 Htb1 Kap114 134: Dpb4 Isw2 Itc1	Abf2 Act1 Bro1 Chd1 Hhf1 Hmo1 Ktr6 Pho23 Reb1 Rps4a Rrp8 Rsc3 Rsc4 Rsc6 Rsc8 Rvb1 Rvb2 Sin3 Ste23 Tra1 Tup1 Vps1 Ybr095c Yil056w	Nu
69	Pre-mRNA cleavage and polyadenylation factor 1A	Clp1 Pcf11 Rna14 Rna15	97: Rpl13a Vid24 130: Clp1 Pcf11 Rna14 Rna15	Cft1 Fip1 Hta1 Imd2 Imd3 Imd4 Pap1 Rpl6b Rps11a Rps24a Sec31 Ssa3	Cy, Nu
70	Small subunit processome	Nan1 Utp10 Utp15 Utp4 Utp8 Utp9	26: Nan1 Utp10 Utp15 Utp4 Utp7 Utp9 28: Pwp2 Utp18 Utp21 Utp6 38: Emg1 Noc4 Nop14 Utp22 65: Bms1 Rcl1 78: Kre33 Ygr145w 80: Bud21 Imp3 86: Dim1 Rrp12	Dip2 Ecm16 Enp1 Imp4 Kri1 Krr1 Mpp10 Mrd1 Nop1 Nop58 Pno1 Rok1 Rpl1b Rps13 Rps17b Rps24b Rps8a Rrp5 Rrp7 Rrp9 Siki Sof1 Ssb2 Utp20 Utp30 Ylr003c	No
71	Complex 71		17: Has1 Nsa1 Rpf1 48: Fpr4 Puf6 Spb4 87: Nip7 Rpl23a 124: Urb1 Urb2	Arx1 Bms1 Brx1 Cdc60 Cin1 Dbp10 Dbp9 Drs1 Erb1 Loc1 Nan1 Noc3 Nog1 Nop13 Nop2 Nop7 Nug1 Pol5 Pwp1 Rix7 Rpl11a Rpl15b Rpl1a Rpl20b Rpl23b Rpl27b Rpl7b Rpl8b Rpp2b Rps22b Spb1 Ssb2 Utp8 Ypl207w Ytm1	Cy, No, Nu
72	Complex 72	Bzz1 Ecm25	127: Edel1 Syp1	Rpl11a Rpl12a Rpl13b Rpl16b Rpl19b Rpl2a Rpl35a Rpl38 Rpl6b Rps11a Rps14b Rps6a Stm1	Cy
73	RNA polymerase II complex	Iwr1 Rpb11 Rpb2 Rpb3 Rpb4 Rpb7 Rpb9 Tfg1 Tfg2	30: Iwr1 Rpb11 Rpb2 53: Spt4 Spt5 76: Rpo21 Yer139c 121: Rpa12 Rpa135 Rpa190 Rpa43 Rpb5 Rpe40	Aos1 Asr1 Med6 Met10 Rpa49 Rpb10 Rpb8 Rpo26 Taf14	Nu
74	mRNA guanylyl transferase complex	Ceg1 Cet1	15: Skp1 Ymr258c Ypt52 18: Ppz1 Ykl088w 19: Lsm12 Pbp1 104: Mrs6 Pyk2	Aro1 Cet8 Ckb2 Fun12 Gdh1 Nip1 Spt16 Ssa3 Ssa4 Top2 Vps3	Cy, Nu
75	19/22S regulator	Ecm29 Rpn1 Rpn10 Rpn11 Rpn12 Rpn13 Rpn5 Rpn6 Rpn7 Rpn8 Rpn9 Rpt1	57: Pre4 Pre8 93: Nas6 Rpn3 141: Pre10 Pre2 Pre3 Pre6 Pup3 Scl1	Blm3 Cdc6 Mlh2 Pre1 Pre5 Pre7 Pre9 Rad50 Rpn2 Rps18b Rpt2 Rpt3 Rpt4 Rpt5 Rpt6 Sro7 Ubp6 Yol087c Yol098c	Cy, Nu
76	Complex 76	Acs2	60: Acs2 Rnr1 89: Ald6 Idh2 108: Sam1 Sam2 109: Rpl8a Yer036c	Cct4 Cct6 Cdc19 Dbp8 Efb1 Eft2 Gpm1 Gsh1 Gsp2 Hsp60 Ilv1 Mae1 New1 Pet9 Rpl3 Rpn1 Rps3 Sec27 Tef2 Tef4 Tsa1 Wrs1 Ybr025c Yef3	Cy
77	Polyadenylation Factor I	Cft1 Cft2 Fip1 Mpe1 Pap1 Pfs2 Pta1 Ysh1 Yth1	23: Ade16 Glc7 Sol2 97: Rpl13a Vid24 130: Clp1 Pcf11 Rna14 Rna15	Bud14 Gsy2 Hca4 Hta1 Imd2 Pti1 Ref2 Rpl16b Rpl6b Rpl8a Rsa3 Sds22 Sec31 Ssa3 Ssu72 Swd2 Syc1 Tk11	Nu
78	Ppz1 protein phosphate complex	Ppz1 Ykl088w	18: Ppz1 Ykl088w	Cka2 Ckb2 Sds22 Ssa3	Cy, Nu
79	Protein phosphatase 1 complex	Ade16 Glc7 Sol2	23: Ade16 Glc7 Sol2	Bmh1 Bud14 Cft1 Gsy2 Pap1 Pti1 Ref2 Reg1 Rnr1 Sds22 Ste23 Swd2 Ysh1	Cy, Nu
80	Prp19-associated complex	Cdc40 Cef1 Clf1 Cwc2 Cwc22 Ecm2 Prp19 Prp45 Prp46 Syf1	1: Lsm6 Lsm7 Prp4 Prp6 Prp8 Snu114 7: Mud1 Nam8 Prp40 Prp42 Snp1 Snu56 Snu71 Sto1 Yhe1 9: Cwc23 Prp43 Ylr424w 13: Prp3 Prp31 Smb1 Spp381 83: Prp39 Smd1 106: Luc7 Smd3 142: Cus1 Hsh155 Lea1 Prp21 Rse1 145: Lsm1 Lsm2 Lsm4 Lsm5 Pat1	Aar2 Brr2 Bud31 Cbc2 Csn12 Dhh1 Dib1 Hsh49 Isy1 Lin1 Msl1 Prp11 Prp22 Prp24 Prp38 Prp9 Seh1 Smd2 Snt309 Snu23 Snu66 Ybl104c Ykr022c Ynl224c	Cy, Nu
81	SAGA complex	Ada2 Hfi1 Ngg1 Sgf29 Sgf73 Spt20 Spt3 Spt7 Spt8	84: Gcn5 Taf10 85: Med6 Yap1 146: Taf1 Taf11 Taf12 Taf3 Taf4 Taf9	Arp4 Epl1 Esa1 Hem13 Med4 Nam7 Nut1 Pdr1 Pgd1 Psk1 Sec2 Sin4 Snf12 Spt15 Taf5 Taf6 Taf7 Tao3 Tra1 Ubp8 Vas1 Vid21	Nu
82	Karyopherins complex	Kap95 Nup2 Nup60 Srp1	14: Pse1 Ulp1 64: Kap123 Ypr174c 107: Asc1 Ynl224c	Fun19 Idh2 Ilv5 Ilv6 Nup1 Pop1 Pop3 Pop8 Rfa3 Rpl18a Rps18a Rps1b Rvb2 Slx9 Spa2 Ura2 Ymr310c Yra1	Cy, NP

83	Sit4/Sap185 protein phosphatase complex	Cdc25 Hrk1 Mds3 Sap185 Sit4	51: <i>Cdc25 Mds3</i>	Acc1 Bem2 Cct2 Cct5 Dig1 Fab1 Mrp13 Pph21 Rpl11b Rts3 Sap155 Sap190 Yef3 Ykr027w	Cy, Nu
84	Nup84 sub-complex	Nup120 Nup145 Nup84 Nup85 Sec13 Seh1	11: Sec16 Sec23 Sfb2 71: Grh1 Sec24	Cys4 Ero1 Mex67 Sec31 Stt3	NP
85	Translational release factor complex	Sup35 Sup45	99: Sen54 Tuf1	Crm1 Mec1 Mrp123 Nab6 Pwp1 Rom2 Rpl16b Rpl36b Rps4b Rps7a Ste23 Sui3 Tao3 Uso1 Vac14 Vps5	Cy, Va
86	Pyruvate dehydrogenase	Lat1 Pda1 Pdb1 Pdx1	14: Pse1 Ulp1 140: Lpd1 <i>Pdb1 Pdx1</i> 143: Kgd1 Kgd2 Ymr31	Cym1 Doa4 Dur1,2 Hsp42 Mss51 Prp46 Spa2	Mt
87	Complex 87	Cdc55 Iml1		Ecm5 Yfr006w	Cy
88	Complex 88	Gaa1 Pgi1			
89	Complex 89	Atg3 Mtr10		Cdc19 Tub1	Cy, Nu
90	Complex 90	Mam33 Mrps35	73: <i>Mrps35</i> Nam9	Bni1	Mt
91	Complex 91	Psa1		Acc1 Cct8 Rps14a Rps24a Rpt1 Sec27 Ssk2 Tef4 Tsa1 Vas1 Yol098c	Cy
92	COPI complex	Cop1 Ret2 Ret3 Sec21 Sec26 Sec27 Sec28	42: Erv41 Erv46 118: Mrp139 Mrp144 139: <i>Cop1 Ret2 Ret3 Sec26 Sec27 Sec28</i>	Ero1 Fks1 Gpt2 Img2 Mrp110 Mrp117 Mrp124 Mrp127 Mrp13 Mrp136 Mrp16 Pet56 Rot2 Tdh1 Utp20	Go
93	Complex 93	Ubp2 Yor138c		Rsp5 Spb1	Cy
94	Complex 94	Hmo1		Arp9 Hhf2 Mot1 Net1 Rpp0 Rps17a Rsc8 Rtt102 Sth1 Vps1	No, Nu
95	Actin capping complex	Cap1 Cap2 Yer071c Yir003w	135: <i>Cap1 Cap2 Yir003w</i>		Cs
96	Gyp5/Gyl1 complex	Dbp2 Gyl1	8: <i>Gyl1</i> Gyp5	Cdc46 Rps0b Rpt6	Bd, Cy, Nu
97	Complex 97	Ydl033c Ygl039w			
98	Complex 98	Lhp1 Sxm1 Utp30			Nu
99	Complex 99		32: Tbf1 Vid22 66: Hta2 Htb1 Kap114 134: Dpb4 Isw2 Itc1	Abf2 Act1 Arp8 Bro1 Hmo1 Lhp1 Mot1 Nap1 Pdr1 Reb1 Rfc1 Rpo31 Rps4a Rps4b Rrp8 Rsc2 Rsc30 Ste23 Tfc4 Yil056w	Cy, Nu
100	Yeats nexin complex	Snx4 Snx41		Atg20 Sla2	Va
101	Complex 101	Dbp8 Rps16a			
102	Complex 102	Npl3 Nrp1		Rps24b	Cy
103	Complex 103	Sgn1	35: Cdc33 Tif4632 115: <i>Sgn1</i> Ygr250c	Boi1 Cyr1 Nab6 Nrd1 Pab1 Sen1 Srv2 Tdh3	Cy
104	Complex 104	Vma10 Vma2	75: <i>Vma2</i> Vma5	Rav1 Tfp1 Vma4 Vma6	Va
105	Complex 105	Hca4 Nrp1		Iml1 Rps4b	Cy
106	Complex 106	Vac8		Bni1 Fba1 Pdc1 Pkg1 Rps17a Tcb3 Vph1 Vtc4 Yor1	Cy, Va
107	Complex 107	Lsm12 Pbp1	19: <i>Lsm12 Pbp1</i>	Eno2 Gln1 Hsp42 Rpp0 Ssa3 Top2	Cy, Nu
108	Complex 108	Pep4 Rtn1			
109	Complex 109	Rpt3 Tub3			
110	Complex 110	Nab6 Nrp1		Npl3 Sgn1 Tif4631	Cy
111	Mannan Polymerase II	Anp1 Hoc1 Mnn10 Mnn11 Mnn9	125: Afg3 Phb1 Phb2 Yta12	Fks1 Gea2 Pmr1 Rpl4b Spo14	Go
112	Complex 112	Lys9 Yor164c		Emi2 Ssa1	Cy
113	SNARE complex	Dsl1 Sec39 Tip20 Ufe1 Use1	46: Pep12 Ykt6	Adh1 Cdc6 Fks1 Sec22 Ths1 Vti1	ER
114	Golgi associated retrograde protein complex	Kar2 Vps52 Vps53 Vps54		Scj1	Go, Va
115	Complex 115	Gln4 Ssl2 Yor352w			Cy, Nu
116	Complex 116	Fun19 Rps5			
117	Complex 117	Rpl24a Rps0a		Clu1 Ded81 Mpm1 Rpl10 Rps17b Rps25b Tim44 Tom40	Cy
118	Complex 118	Aco1 Faa4			
119	Complex 119	Nab6 Pab1			Cy
120	Exocyst complex	Exo84 Sec10 Sec3 Sec5 Sec6 Sec8		Exo70 Rpc17 Sec15 Vps9	Bd, CP
121	Complex 121	Gon7 Kae1 Yml036w			
122	Complex 122	Aco1 Mdh1 Yhb1	110: Hsp60 Mkt1 114: <i>Aco1 Yhb1</i>	Ape2 Bud4 Rad51	Cy, Mt
123	Complex 123	Pro1		Ecm5 Hhf1 Iml1 Psd2 Rpl16a Rpl38 Rps9a Uso1 Yfr006w	Cy
124	Complex 124	Pdr5 Usa1			

125	Complex 125	Rpl33a Rpl35b		Brx1 Dep1 Ebp2 Gen5 Hir2 Pdr1 Rpl12a Rpl1b Rpl23b Rpl25 Rpl27a Rps14b Rrp1 Top1 Ybr095c	Cy, Nu
126	Complex 126	Fpr4 Rix7 Rpl20a		Dbp9 Drs1 Ebp2 Nog1 Puf6 Rpl3 Rpl7b Rps18a Spb1 Yer139c	Cy, No, Nu
127	Nucleotide excision repair factor 3 complex	Ssl1 Tfb1 Tfb4		Cdc27 Fap1 Imd3 Krs1 Mrps28 Tfa1	Nu
128	Complex 128	Ebp2 Yer139c			
129	Complex 129	Tef4	56: Mft1 Sub2	Aap1' Efb1 Eft2 Hpr1 Hrb1 Mtr10 New1 Nmd5 Pfk1 Rlr1 Rpn2 Top2 Usa1 Ybr025c	Cy, Nu
130	Complex 130	Atp3 Cdc37		Rpl10 Sui2	Cy
131	Complex 131	Imp4 Rpn2		Rps3 Rps6a Utp13 Ynk1	Cy
132	Complex 132	Tub3		Atp2 Clu1 Hsp42 Kap122 Tub2 Ydj1	Cy, Nu
133	Complex 133	Aep1 Hrp1 Rps9a		Npl3 Nrp1 Rps11a Rps24b Rps8b	Cy
134	COP9 signalosome complex	Csi1 Csn9 Pci8 Rri1 Rri2		Rpn5	Cy
135	Complex 135	Ilv1 Psa1	55: Cpa1 Cpa2 101: Sse1 Tsa1	Acc1 Cln2 Eft1 Fas1 Hsp60 Rpl18b Rpl36b Rpn2 Rps14a Rpt3 Tub3 Vas1 Yol098c	Cy, Mt
136	Complex 136	Cse1 Sec7	92: <i>Cse1 Sec7</i>	Dhh1 Fks1 Idh2 Ktr3 Mir1 Rpl4a Rpp0 Yfr044c	Cy, Mt
137	Act1/Sac6 complex	Act1 Sac6		Mgs1	
138	Complex 138	Rpl31a Rpl33b		Rps18b Rps24a	Cy
139	Complex 139	Rli1		Atg9 Dbp2 Fun12 Gcd11 Hcr1 Imd3 Nat1 Nip1 Rpg1 Rps11a Rps11b Rps17a Rps18a Rps22a Rps22b Rps4b Rps6a Rps7a Rps8b Rps9a Sui2 Sui3 Tif34 Tif35 Tif5	Cy
140	Complex 140	Atg9 Rpl18b Rpl7a		Ipi1 Nog1	Cy, Nu
141	Complex 141	Cam1 Hsl1 Ydr314c		Efb1	
142	Ctf19 protein complex	Chl4 Ctf19 Ctf3 Mcm21 Mcm22 Okp1	27: Dsn1 Mtw1	Ame1 Nkp1 Nsl1	Nu
143	AP-2 complex	Apl1 Apm4		Ltv1 Rpl16a Rpl17a Rpl1b Rpl2b Rpl35a Rpl8b Rps17a Rps1a Rps6a	Bd, CP, Cy
144	Complex 144	Faa4 Ypl206c		Aco1 Clu1	Va
145	Chaperonine containing T-complex TRiC	Cct2 Cct7 Plp2 Vid27		Cct4 Cct5 Cct6 Cct8 Hgh1 Ilv6 Tub2	
146	RNA polymerase II general transcription factor TFIID	Taf2 Taf8	21: Taf14 <i>Taf2</i> Taf8	Taf1 Taf10 Taf5 Taf6 Taf7 Tef4	Nu
147	Complex 147	Bbc1 Yjl123c	119: Ids2 Pep4	Tra1 Yhr029c	Cy
148	PAN complex	Pan2 Pan3			Cy
149	Complex 149		28: Pwp2 Utp18 Utp21 Utp6 38: Emg1 Noc4 Nop14 Utp22 62: Imp4 Rrp9 Utp13 65: Bms1 Rcl1	Bfr2 Cka1 Cka2 Ckb1 Ckb2 Dim1 Dip2 Ecm16 Fas2 Gbp2 Imp3 Mpp10 Mrd1 Nop1 Nop58 Pno1 Rok1 Rpl26b Rps11a Rps13 Rps16a Rps17a Rps1b Rps24b Rps4a Rps4b Rrp7 Utp10 Utp30 Utp7 Ygr145w Ylr003c Yor059c	Cy, No, Nu
150	Complex 150	Atg9 Rps6a		Rps3 Rps7b	Cy
151	Complex 151	Dig1 Npl3 Pep1		Sap190	Nu
152	Bre5/Ubp3 complex	Bre5 Ubp3	44: <i>Bre5 Ubp3</i>	Cdc19 Idp1 Nop1 Rad50 Rpl1b Rps1a Tor1	Cy
153	Tid3 complex	Nuf2 Spe24 Spe25 Tid3			Nu
154	Complex 154	Dph2 Eft1 Yil103w			Cy
155	Complex 155	Atg11 Atg17 Ypl166w			Va
156	Complex 156	Met10 Rpa135		Aha1 Dnm1 Gen1 Gfa1 Lys12 Rpc19	Cy, No
157	Complex 157	Esf1 Krr1 Mvd1 Nop6 Rpl26b Ygr145w	80: Bud21 Imp3	Bfr2 Bms1 Cbf5 Dbp8 Ero1 Rpl15a Rps16a Rps18a	Cy, No
158	Complex 158	Aos1 Spt4	53: <i>Spt4</i> Spt5	Rpb11 Rpb4 Rpb5 Rpb7 Rpo26 Tfg1 Uba2 Yef3	Nu
159	Complex 159	Cdc60 Nan1 Utp15 Utp4	86: Dim1 Rrp12	Pno1 Pol5 Rpl1b Utp10 Utp8 Utp9	No
160	Complex 160	Rpl33b Rpp2b Rsa3		Mak5 Rpl16a Rpl18a Rpl18b Rpl2a Rpl31a Rpl36b Rpl43a Rpl6a Rps16b Rps18b Yor1	Cy
161	THO complex	Hpr1 Hrb1 Mft1 Rlr1 Sub2 Tex1 Thp2	56: <i>Mft1 Sub2</i>	Atp2 Gbp2 Imd3 Nop1 Rpn2 Rps4a Tef4 Top2 Ybr025c Yra1	Nu
162	Complex 162	Rfc1 Rfc2		Imd3 Rfc3 Rps0b	Nu
163	Ccr4/Not protein complex	Caf40 Ccr4 Cdc39 Pop2	82: Pol1 Pol12 Pri1 Pri2	Atg9 Caf130 Rps6b Rvb1 Rvb2	Cy
164	Complex 164	Mis1 Pim1		Rpl10 Rpl17a Rpl1b Rpl27b Rpl6b Rps1b	Cy, Mt

165	Complex 165	Tub3	25: Erp3 Erp5 63: Mae1 Tom1 81: Gcn1 Gen20 108: Sam1 Sam2	Acc1 Cct4 Cct6 Clu1 Crm1 Fks1 Kap122 Mdn1 Msn5 Por1 Psa1 Rpl1a Rpl4a Rpn1 Rpn2 Rpt3 Rpt4 Rpt6 Ths1 Ura7	Cy, ER, Nu
166	Complex 166	Clu1 Kap104		Mtr10 Nmd5 Usa1	Cy, Nu
167	Svg1/cyclin Bur2 protein complex	Bur2 Sgv1	123: <i>Bur2 Sgv1</i>	Brr1 Hta1 Mud1 Nrd1 Scp160 Sen1 Smd3 Sro9 Ssb2	Nu
168	Complex 168	Pfk2 Rps10a Rps22a			No, Nu
169	Complex 169	Ade3 Cse1 Tdh1 Tom1 Ydr341c		Aap1' Fas1 Fas2 Fur1 Grs1 Hsp82 Ils1 Ilv5 Ilv6 Msn5 Sar1 Sec53 Tdh3 Trm3 Vps13 Ynk1	Cy, Nu
170	Complex 170	Hmg1 Rad51 Rpl27b			Cy
171	Complex 171	Gcn1	89: Ald6 Idh2 116: Hsp82 Lys12	Aha1 Crm1 Cse1 Gfa1 Kap123 Mam33 Met10 Mpm1 Msn5 Pdc1 Pet9 Ptm1 Rps0b Sam1 Sec53 Ssa4 Tub1	Cy, Mt, Nu
172	Complex 172	Ded81		Clu1 Lys12 Mdn1 New1 Rpl10 Rpl24a Sam1 Ufd4 Yhr020w	Cy
173	Complex 173	Nam9	59: Mrp1 Rsm25 73: Mrps35 <i>Nam9</i>	Ade6 Air2 Ehd3 Met6 Mrp13 Mrp17 Mrp21 Mrp4 Mrp51 Mrps16 Mrps17 Mrps18 Mrps28 Mrps5 Mrps8 Mrps9 Pet123 Psd2 Pus7 Rsm10 Rsm22 Rsm23 Rsm24 Rsm26 Rsm7 Trf4 Ubp10	Mt
174	Complex 174	Yer036c	109: Rpl8a <i>Yer036c</i>	Dbp8 Mlp1 Rpl15a Rpl20a Rpl3 Rpl6a Rps11b Vma13	Cy
175	Complex 175	Brr2 Cus1 Lea1 Prp21 Prp3 Prp31 Prp4 Prp6 Prp8 Prp9 Smb1 Snu114 Snu23 Snu66 Spp381	1: Lsm6 Lsm7 <i>Prp4 Prp6 Prp8 Snu114</i> 13: <i>Prp3 Prp31 Smb1 Spp381</i> 83: Prp39 Smd1 142: <i>Cus1 Hsh155 Lea1 Prp21 Rse1</i> 145: Lsm1 Lsm2 Lsm4 Lsm5 Pat1	Aar2 Bud31 Cef1 Clf1 Csn12 Cwc22 Cwc23 Dhh1 Dib1 Ecm2 Hsh49 Isy1 Krs1 Lin1 Mrpl8 Msl1 Prp11 Prp24 Prp38 Prp46 Seh1 Smd2 Smd3 Syf1 Ybl104c Ylr424w	Nu
176	Complex 176	Rpl19a Rps14a Rps6b		Rpl26b	Cy
177	Complex 177	Rpl33b Rps16b Rsa3 Yor1		Dbp9 Dcs2 Nop12 Rpl10 Rpl15a Rpl17b Rpl18b Rpl26b Rpl31a Rpl43a Rpl4b Rpp2b Rps14a Rps18b Rps24a Rps24b Rrp5 Tfb3	Cy
178	Complex 178	Bud3 Sec1		Ded1 Iml1 Mrpl20 Mrpl8 Snf2	Bd
179	Ku complex	Yku70 Yku80	136: <i>Yku70 Yku80</i>		Nu
180	Complex 180	Rpn2 Yol098c			Cy
181	Rubiquitin -activating complex	Uba3 Ula1			
182	TRAPPII complex	Bet3 Bet5 Gsg1 Kre11 Trs120 Trs130 Trs20 Trs23 Trs31 Trs33		Fks1 Skn1	Go
183	Complex 183	Ade5,7 Rps5		Rps3	
184	Kel1/Lte1 complex	Bem2 Kel1 Lte1	12: <i>Kel1 Lte1</i>	Fks1	Bd, Cy
185	Complex 185	Ald6 Ptc2 Shq1	36: Ptc3 Ydr071c		Cy, Nu
186	Complex 186	Gln1 Hsp42		Mdn1 Rpb2 Ura2	Cy
187	Complex 187	Brr1 Hta1			
188	Origin of replication recognition complex	Orc1 Orc2 Orc3 Orc4 Orc5 Orc6		Pab1 Rpl16b	Cy, Nu
189	Complex 189	Prp43 Ynl224c	107: <i>Asc1 Ynl224c</i>	Nop1 Prp46 Rpl12b Rpl31b Rps10b Rps16b Rps20 Rps22a Rps4a Rps8a Rps9a Sbp1 Scp160 Sro9 Ylr424w	Cy, No, Nu
190	INO80 protein complex	Arp5 Arp8 Ies1 Ies3 Ies5 Ino80 Nhp10	54: <i>Arp5 Rvb1</i> 100: Rvb2 Sec53	Abf2 Act1 Arf1 Arp4 Atp1 Hsp82 Isw2 Itc1 Ktr6 Met10 Pmr1 Rad50 Rpt6 Sar1 Taf14 Vps1	Cy, Nu
191	Complex 191	Myo5	68: Cmd1 <i>Myo5</i>	Cdc47 Crm1 Eht1 Osh2 Pol5 Rps0b	Cs
192	Complex 192	Ptc4 Ydr186c	6: <i>Ptc4 Ydr186c</i>	Ssb1	Cy
193	Complex 193	Eno1 Nat1		Asc1 Rpl31a Rpl7b	Cy
194	Complex 194	Yor1		Atp1 Erb1 Mak5 Rpl10 Rpl18b Rpl27b Rpl30 Rpl33b Rpl36b Rpl4a Rpl7b Rps18b Rps1a Rsa3 Vma10 Vph1 Ytm1	Cy, No, Nu, Va
195	Set3 histone deacetylase protein complex	Cpr1 Hos2 Hos4 Set3 Sif2 Snt1		Kap95 Pph22 Rim1 Snf2 Zds1	Cy, Nu
196	Complex 196	Rpl19a Rpl9a			Cy
197	Complex 197	Abf2 Vps1		Pdr1 Reb1 Rsc1 Rsc2 Rsc3 Sth1 Top1 Yil056w	Cy, Nu
198	Complex 198	Ebp2 Rpl23b	17: Has1 Nsa1 Rpf1 52: Cic1 Ipi1 61: Noc2 Nop12 Rrp5 87: Nip7 Rpl23a	Brx1 Dbp10 Mrt4 Noc3 Nop15 Rix1 Rix7 Rpl12a Rpl14a Rpl20b Rpl31b Rpl35b Rpl5 Rpl7b Rrp1 Spb1 Spb4 Ytm1	Cy, No, Nu

199	Complex 199		65: Bms1 Rcl1 69: Ecm16 Krr1 Nop6	Emg1 Enp1 Gbp2 Kre33 Kri1 Noc4 Nop1 Nop14 Rpb10 Rps11b Rps17a Rps20 Rps4b Rps8a Rps8b Rrp12 Sof1 Utp10 Utp20	Cy, No, Nu
200	Complex 200	Dur1,2 Mrpl22 Pro1		Acc1	Cy
201	RNase MRP	Pop1 Pop3 Pop4 Pop5 Pop8 Rpp1			Nu
202	Complex 202	Sui1		Prt1 Rps10a Rps22a	No, Nu
203	Complex 203	Gbp2 Thp2		Hpr1 Imd3 Nop1 Rpl8b Rps4b Sub2	Cy, Nu
204	Complex 204	Hem13 Psk1 Vas1			Cy
205	Complex 205	Pep4 Ymr196w	119: Ids2 <i>Pep4</i>	Rtn1	Cy
206	Complex 206	Mrp17 Ubp10	59: Mrp1 Rsm25	Bem2 Ecm16 Ehd3 Mrp13 Mrp21 Mrp4 Mrp51 Mrps18 Mrps28 Mrps5 Pho23 Psd2 Pus7 Rsm10 Rsm19 Rsm22 Rsm23 Rsm26 Rsm27 Utp20	Mt, Nu
207	Complex 207	Aar2 Bud31 Lin1		Snt309	Nu
208	Complex 208	Ids2 Pep4	119: Ids2 <i>Pep4</i>		
209	DNA polymearse alpha-primase complex	Pol1 Pol12 Pri1 Pri2	82: <i>Pol1 Pol12 Pri1 Pri2</i> 126: Rfc1 Rfc3 Rfc5 144: Srp101 Srp102	Rfc2 Rps6b Uso1	Nu
210	Complex 210	Dbp9 Spa2 Ymr310c		Pre8	Nu
211	Complex 211	Bud21 Rrp9 Sik1	28: Pwp2 Utp18 Utp21 Utp6 38: Emg1 Noc4 Nop14 Utp22 62: Imp4 <i>Rrp9</i> Utp13 65: Bms1 Rcl1 72: Hca4 Nop58 <i>Sik1</i> 80: <i>Bud21</i> Imp3	Bfr2 Dbp8 Dim1 Dip2 Ecm16 Enp1 Ero1 Esf1 Kre33 Kri1 Mpp10 Mvd1 Nop1 Nop6 Pno1 Pre9 Rok1 Rpl19a Rpl26b Rpl33a Rps16a Rps1a Rps4b Rps8b Rrp5 Sof1 Utp10 Utp20 Utp7 Utp9	No
212	Complex 212	Tpi1		Eno2 Far3 Fba1 Gpm1 Nup1 Pkg1 Rhr2 Sse2 Ydr161w	Cy, Nu
213	Complex 213	Ade5,7 Reb1 Top1		Npl6 Rps3 Rps4b Rps5 Rsc2 Rsc3	Cy, Nu
214	Complex 214	Ktr3 Rpl4a			
215	Complex 215	Ade6 Rsm23			
216	Complex 216	Pse1 Slx9 Spa2 Ulp1	14: <i>Pse1 Ulp1</i> 57: Pre4 Pre8	Dbp9 Kap95 Mds3 Pre5 Rfa3 Rpt5 Rrp12 Scl1 Srp1 Ymr310c Yol054w Yra1	Cy, NP, Nu
217	Complex 217	Aro3 Ilv5			
218	C-terminal domain kinase I complex	Ctk1 Ctk2 Ctk3		Rpl1b Rpl20a Rps13 Rps9a Ssa1 Sse2 Yor164c	Cy, Nu
219	Spindle pole body	Pfk1 Spc72 Spe97 Spe98 Tub4			Nu
220	Complex 220	Psa1 Yol098c	101: Sse1 Tsa1 137: Smc1 Smc3	Aro1 Cct8 Cdc19 Ilv1 Rpn2 Rps0b Rps14a Rps24a Rpt1 Vas1	Cy
221	Complex 221	Imd4 Tdh1		Cdc33 Imd2 Kar2 Rpl1b Rps8b Tdh2	Cy, Nu
222	Complex 222	End3 Lsb3 Sla2			Va
223	Complex 223	Sen54 Tuf1	99: <i>Sen54 Tuf1</i>		Mt
224	SWI-SNF global transcription activator complex	Arp9 Snf11 Snf12 Snf5 Snf6 Swi1 Swi3	24: <i>Arp9 Swi1 Yfl049w</i> 147: <i>Snf11 Snf2 Snf5 Snf6 Swi3</i>	Arp7 Pdr1 Rtt102 Taf14 Vps1 Yap1	Nu
225	Complex 225	Ymr310c	98: Cbf5 <i>Ymr310c</i>	Aar2 Dbp2 Hek2 Mis1 Mrp7 Nab6 Nop1 Rpl13b Rpl43a Rpl6b Rps17a Rps24a Rps4a Rps6a Rrp12 Slx9 Snu66 Spa2 Sro9 Stm1 Sui3	Cy, Nu
226	Boi2/Bem1/Cdc24 complex	Boi2 Cdc24		Cdc123 Rpl12b Rpl17b Rpl4b Rpp0 Vid24	Cy, Nu
227	Rai1/Rat1 complex	Rai1 Rat1	20: Vps29 Vps35 120: <i>Rai1 Rat1</i>	Bem2 Cdc25 Cdc48 Mrps28 Pfk2 Rpl11a Rpl20b Rpl8b Rpp0 Rps10a Rps14b Rps17b Rps20 Rps22a Shp1 Yal036c Ydr049w Ypl009c	Cy, Nu
228	Complex 228	Ipp1 Rom2 Ydl203c		Mex67 Nup85 Rpl13b Rpl16a Rpl27a Rpl2b Rps14b Tus1 Ypl066w	Cy, NP
229	Complex 229	Eft2 Hgh1			Cy
230	Complex 230	Ecm30 Ubp15		Ded1 Doa4 Tub2 Ura2	Cy
231	Complex 231	Mrpl36 Pet56			Mt
232	DNA polymerase delta	Cdc2 Hys2			Nu
233	Anthranilate synthase complex	Sam1 Trp2 Trp3	34: <i>Trp2 Trp3</i> 81: Gcn1 Gcn20	Aha1 Aro1 Dnm1 Hsc82 Met10 Rpa135 Rpn1 Rpt3 Tub3 Ufd4	Cy
234	Complex 234		10: Rgp1 Ric1 65: Bms1 Rcl1 79: Nob1 Rps18b 86: Dim1 Rrp12	Ded1 Dip2 Glt1 Mpp10 Mrd1 Pno1 Prt1 Rio2 Rpg1 Rps0a Rps10a Rps11a Rps13 Rps21a Rps22a Rps28b Rps3 Rps5 Rps7b Rps8a Rps9b Sui1 Tif34 Tif5 Utp20 Ygr054w	Cy, Go
235	SBF, SWI4-SWI6- dependent cell cycle box binding factor complex	Swi4 Swi6		Tdh3	Cy, Nu

236	Pms1/Mlh1 mismatch DNA repair complex	Mlh1 Pms1		Mlh2 Ssa2	Cy, Nu
237	Complex 237	Nrp1	74: Cbc2 Npl3 115: Sgn1 Ygr250c	Hca4 Hrr25 Iml1 Kem1 Lhp1 Mtr4 Nab3 Nab6 Nam7 Nam8 Nrd1 Pab1 Rio2 Rps11a Rps22a Rps8b Rps9a Sen1 Snp1 Snu56 Ste23 Tef1 Tif4631 Tif4632	Cy, Nu, Va
238	Complex 238	Dbp9 Spa2			
239	Small subunit processome	Dip2 Mrd1 Nan1 Nop58 Pno1 Pwp2 Rok1 Rrp9 Utp10 Utp13 Utp15 Utp18 Utp21 Utp22 Utp4 Utp6 Utp7 Utp8 Utp9 Ylr003c	26: Nan1 Utp10 Utp15 Utp4 Utp7 Utp9 28: Pwp2 Utp18 Utp21 Utp6 62: Imp4 Rrp9 Utp13 80: Bud21 Imp3	Cdc60 Dim1 Mpp10 Nop1 Rps13 Rrp5 Sik1 Sof1 Yor059c	No
240	Complex 240	Tif4631 Yal036c	3: Dpb2 Pol2	Cbf5 Cdc33 Gbp2 Imd4 Kem1 Lhp1 Mrp7 Pus7 Rpl31b Rps10b Rps20 Rps22a Rps6a Rrp8 Sbp1 Sgn1 Snp1 Sro9 Sto1 Tif2	Cy, Nu
241	Complex 241	Kex2 Sil1 Ybt1			Va
242	Complex 242	Adh3 Nam7			
243	Phosphoribosyl diphosphate synthase complex	Prs1 Prs3		Eft1 Rpl30	Cy
244	Complex 244	Ykr018c		Tfa1 Ygr130c Ynr029c	Cy, Nu
245	Complex 245	Dbp8 Rpl15a		Rpl16b	
246	Complex 246	Dhh1 Sec7	92: Cse1 Sec7	Ktr3 Yfr044c	Cy
247	Complex 247	Aro1 Sam1			Cy
248	Protein phosphatase 2A complex	Cdc55 Rts3 Tpd3 Zds1 Zds2	12: Kell1 Lte1	Cin1 Iml1 Pph21 Pph22 Rps17a Rps1b Rps24a Rps6a Rts1 Vph1 Ybl104c Ykr027w Yor1	Cy, Nu
249	Translation elongation factor EF1	Cam1 Efb1 Tef2		Eft2 Hsl1 New1 Rpn1 Rps24b Sbp1 Tef4 Ybr025c Ydr314c	Cy
250	Complex 250	Bms1 Mrd1	65: Bms1 Rcl1	Ecm16 Rps7b Utp7 Ygr054w	
251	Complex 251	Pdr16		Aap1' Acc1 Dim1 Rps11b Rps17a Rrp7 Tra1 Yef3 Yra1	Nu, Va
252	Complex 252	Rpl13a Rpl2a	97: Rpl13a Vid24 113: Fun12 Rpl12b	Atg9 Lsg1 Mak5 Mis1 Nip1 Rpl10 Rpl11b Rpl13b Rpl15a Rpl16b Rpl18b Rpl20a Rpl24a Rpl25 Rpl26b Rpl27a Rpl27b Rpl2b Rpl33a Rpl33b Rpl36a Rpl36b Rpl6a Rpp0 Rpp2b Rps14a Rps14b Shm1 Yor1 Ypl009c	Cy
253	Complex 253	Yil056w		Abf2 Act1 Arp8 Hhf1 Hmo1 Htb1 Ies1 Ioc3 Isw2 Itc1 Mot1 Nhp10 Reb1 Rfa1 Rpe37 Rsc1 Rsc2 Rsc3 Rsc30 Sth1 Tfc4 Vps1	Cy, Nu
254	Complex 254	Ade3 Dps1 Sar1 Tdh1 Ydr341c Ynk1	63: Mae1 Tom1 81: Gcn1 Gcn20 100: Rvb2 Sec53	Aap1' Cct4 Cse1 Eft2 Fas1 Fas2 Fur1 Grs1 Hhf2 Hsp82 Ils1 Ilv5 Ilv6 Msn5 Pfk2 Rpa135 Rpl30 Rps3 Rpt5 Sam1 Ssk2 Tfp1 Ths1 Trm3 Tub2 Ura2 Vps13	Cy, Nu
255	Complex 255	Rok1 Rps1b		Nop6 Rpl20a Rps17a Rps24a Rps8b	Cy
256	Chromatin assembly complex	Cac2 Msi1			Nu
257	Complex 257	Mpm1 Rps25b Tom40 Ygr130c	94: Kap104 Mir1	Clu1 Ecm29 Rpl24a Rps9a	Cy
258	Kap104/Hrp1 complex	Hrp1 Kap104		Clu1 Rpl12a Rpp0 Rps13 Rps8b	Cy, Nu
259	Complex 259	Fas1 Trm3			Cy
260	Complex 260	Dcs2 Tfb3	109: Rpl8a Yer036c	Far3 Ist2 Rpl11b Rpl14b Rpl17a Rpl17b Rpl25 Rpl26b Rpl28 Rpl2a Rpl2b Rpl3 Rpl33b Rpl35b Rpl6a Rps14a Rps16a Rps18a	Cy
261	Complex 261		30: Iwr1 Rpb11 Rpb2 76: Rpo21 Yer139c	Asr1 Dbp10 Dps1 Mak21 Nsa2 Nug1 Rpb3 Rpb4 Rpb5 Rpb7 Rpb8 Spb1 Ura2	Cy, Nu
262	Clathrin-associated protein AP-3 complex	Apl5 Apl6 Apm3		Eno2	Go, Va
263	Complex 263	Aar2 Gcn2 Snu66			Cy, Nu
264	Complex 264	Ade6 Pet123			
265	Complex 265	Fun12 Rpl12b	113: Fun12 Rpl12b	Nip1 Rpl13a Rpl2a Rps22a Ypl009c	Cy
266	Complex 266	Mrd1 Utp22 Utp7 Ylr003c		Dip2 Mpp10 Nop1 Nop58 Pno1 Pwp2 Rok1 Rrp5 Rrp7 Utp20 Utp30 Utp6	No, Nu
267	Complex 267	Imd2 Lys2 Sxm1			Cy
268	Complex 268	Adh1 Prc1			
269	Complex 269	Mec1 Vac14			
270	Complex 270	Clu1 Ura7			Cy
271	alpha-Aminoacidipate-semialdehyde dehydrogenase complex	Lys2 Lys5		Rps4b Sxm1	Cy
272	Complex 272	Asc1 Nat1		Clu1 Eno1 Mis1 Rpl6b Rpl7b	Cy
273	Complex 273	Fap1 Yil161w		Imd3 Rpl17b Rpl2b Rpl4a Rpl5 Rpl7a Rpp0 Rps0a Rps22a Rps22b Rps8b Skt5	Cy

274	Complex 274	Tcb3		Ktr3 Rpl1a Rpl28 Rpl4a Rpl4b Rpp0 Trm3 Ymr134w	Cy
275	Complex 275	Pdc1 Ptm1			
276	Ste50/Ste11 complex	Ste11 Ste50			Cy
277	ATP-dependent Isw2p-Itc1p chromatin remodeling complex			Act1 Arf1 Arp4 Arp8 Atp1 Atp2 Atp3 Gfa1 Ies1 Ies3 Ies5 Ino80 Isw2 Itc1 Nhp10 Rad50 Rvb2 Sar1 Tfp1 Vps1	Mt, Nu
278	Complex 278	Yal036c		Bem2 Cdc25 Kre33 Kri1 Rpl11a Rps10a Rps10b Rps14b Rps20 Rps4a Rps5 Rps7a Scp160 Ssz1 Zuo1	Cy
279	Complex 279	Gdb1 Nup192			
280	Complex 280	Ptc3 Ydr071c	6: Ptc4 Ydr186c 36: Ptc3 Ydr071c	Ptc2 Rad50 Vps15 Wtm1 Yhb1	Cy, Nu
281	Cytoplasmic mRNA processing body	Krs1 Lsm1 Lsm5 Pat1	128: Dcp1 Dcp2 Edc3 145: Lsm1 Lsm2 Lsm4 Lsm5 Pat1	Adh3 Cus1 Dhh1 Hsh155 Kem1 Nam7 Prp11 Prp21 Prp9 Rps8b Rse1 Smd3	Cy, Nu
282	Complex 282	Pfk1 Pfk2	103: Pfk1 Pfk2 108: Sam1 Sam2	Aco1 Aro1 Dnm1 Dps1 Mcm22 Rps10a	Cy
283	Complex 283	Sse2 Yor164c			Cy
284	Complex 284	Cdc36 Mrd1 Rps10b		Not5 Rps13 Rps4a	Cy
285	Complex 285	Asc1 Rpl27b Skt5		Nat1 Rpl4b	Cy
286	Complex 286	Dbp2		Amn1 Caf130 Rpl43a Rps0b Rps7b Ssz1 Stm1 Ygr054w Zuo1	Cy
287	Translation initiation factor eIF2B complex	Gcd1 Gcd11 Gcd2 Gcd6 Gcd7 Gcn3		Ala1 Cct4 Cdc123 Cpa2 Hir2 Ilv1 Met10 Pdr1 Pro1 Rpa135 Rpe19 Rpl11b Rpl12a Rpl12b Rpl33a Rpl35b Rpl36a Rpl36b Sui2 Sui3 Top2 Xks1	Cy
288	Complex 288	End3 Rpc17		Ret1 Rpc19 Rpc25 Rpc34 Rpc37 Rpc53	Nu
289	Complex 289	Chs1 Eno2 Pkg1		Asr1 Fba1 Vac8 Vtc4	Cy
290	Complex 290	Abz1 Mpa43 Tsa1			Cy
291	Complex 291	Rpl35a Rps6b Ura7		Mrpl3 Rpl17b Rpl19a Rpl1a Rpl2b Rpn10 Rpn2 Rps14a Tif1 Tif2	Cy
292	Complex 292	Spo14 Ygl036w			Cy
293	Golgi transport complex	Cog1 Cog2 Cog3 Cog4 Cog5 Cog6 Cog7 Cog8 Im11		Nrp1	Go
294	Complex 294	Fap7		Bfr2 Cbf5 Imp3 Krr1 Mpp10 Nop1 Rps14a Rps14b Rps3 Rps7a Sof1 Utp30 Ygr145w	Cy, No, Nu
295	Complex 295	Pro1 Xks1			Cy
296	Histone acetyltransferase B complex	Hat1 Hat2 Hhf2 Hif1	55: Cpa1 Cpa2 96: Hhf1 Yol054w	Ecm5 Pro1 Psd2 Rpl38 Rps10a Rpt2 Rpt3 Rpt5 Uso1 Xks1 Yfir006w	Cy, Nu
297	Complex 297	Pgi1 Sec1		Kcc4	Bd, Cy
298	Complex 298	Dps1 Ilv1 Rpl18b		Nsa2 Rpb4 Rpl36b Ths1 Tub3	Cy
299	Complex 299	Pab1 Ygr250c			Cy
300	Ufd1/Npl4/Cdc48, ER ubiquitinated proteins degradation complex	Cdc48 Npl4 Shp1 Ufd1 Ydr049w		Bni1 Mrps28 Reg1 Rpl17b	Cy, Nu
301	RAVE/Skp1 complex	Skp1 Ymr258c	15: Skp1 Ymr258c Ypt52 31: Rav1 Rav2 104: Mrs6 Pyk2	Gdh1 Yef3	Cy, Nu
302	Complex 302	Mtf2 Sls1	29: Rvs161 Rvs167	Adh3 Nam7 Rpl27b Skt5	Cy, Mt
303	2-oxoglutarate dehydrogenase complex	Kgd1 Kgd2 Ymr31	79: Nob1 Rps18b 140: Lpd1 Pdb1 Pdx1 143: Kgd1 Kgd2 Ymr31	Cdc19 Rok1 Rpl15b Rpl16b Rpl20b Rpl27b Rpl30 Rpl7a Rps1b Rps24a	Mt
304	Cytochrome bc1 complex	Cor1 Qcr2		Dur1,2 Mrpl22 Npl3 Pro1 Syf1	Cy, Mt
305	Methionyl- and glutamyl-tRNA synthetases complex	Arc1 Gus1 Mes1		Hem13 Vas1	Cy
306	Complex 306	Hua2 Ypl206c		Crm1 Faa4 Yal048c	Va
307	Complex 307	Nop13 Rlp7			
308	Complex 308	Aro3 Wtm1			Nu
309	mRNA cap-binding -Bur2/Sgv1 complexes		35: Cdc33 Tif4632 98: Cbf5 Ymr310c 123: Bur2 Sgv1	Far8 Gsh1 Hek2 Imd2 Imd3 Imd4 Kem1 Kri1 Pab1 Rpl7b Rps10b Rps11b Rps24a Rps24b Rps4a Rps8b Rps9b Tor1	Cy
310	Complex 310	Top1		Abf2 Dep1 Pdr1 Pho23 Rpl17a Rpl35b Rps14b	Cy, No, Nu
311	Complex 311	Fun19 Kap123 Ypr174c	64: Kap123 Ypr174c	Aha1 Gen1 Idh2 Ilv6 Lys12 Nup60 Rpl18a Rps11b Rps18a Rps1a Rps1b Rps5 Tif2 Tub2	Cy, Nu

312	cAMP-dependent protein kinase complex	Bcy1 Tpk2			Nu
313	Complex 313	Rpl18b Tub3		Gfa1 Tub2 Yol098c	Cy
314	Complex 314	Aro1 Cct8	103: Pfk1 Pfk2 114: Aco1 Yhb1	Aha1 Gfa1	Cy
315	AP-1 adaptor complex	Apl2 Apl4 Apm1 Apm2 Aps1		Hek2 Mis1 Pgi1 Rpl11a Rpl16b Rpl19a Rpl19b Rpl1a Rpl3 Rpl36b Rpl6b Rpl9a Rps6a Sda1 Sxm1 Yfl034w	Go
316	Complex 316	Fpr4 Nop13 Puf6 Rix7	17: Has1 Nsa1 Rpf1 48: <i>Fpr4 Puf6</i> Spb4 87: Nip7 Rpl23a 88: Dbp9 Nop4	Arx1 Brx1 Cic1 Dbp10 Drs1 Nog1 Nop2 Nug1 Pol5 Rlp7 Rpf2 Rpl11a Rpl15b Rpl20a Rpl25 Rpl3 Rpl30 Rpl31a Rpl7b Rpl8b Rpp2b Rrp1 Spb1 Ytm1	No, Nu
317	Complex 317	Kar2 Scj1			
318	Complex 318	Dop1 Mon2	2: <i>Dop1 Mon2</i>	Rps0a Ufd4	Cy, Go, Va
319	Complex 319	Gpi17 Ptm1		Pdc1 Tdh3 Tub1	Cy, Nu
320	Ribonucleoside-diphosphate reductase complex	Rnr2 Rnr4	132: <i>Rnr2 Rnr4</i>	Aro3 Dph2 Hir3 Ilv5 Mam33 Rad53 Wtm1 Yil103w	Cy, Nu
321	Complex 321	Erg1 Sec2			
322	Complex 322	Fur1 Rvb2 Sec53	100: <i>Rvb2 Sec53</i>	Ils1 Tub1	Cy, Nu
323	Complex 323	Rpl38 Yol054w	96: Hhf1 <i>Yol054w</i>	Ecm5 Iml1 Rps11b Yfr006w	Cy, Nu
324	Complex 324	Hir3 Mam33 Rad53			Nu
325	Complex 325	Hem13 Taf7	84: Gcn5 Taf10	Ngg1 Psk1 Taf6 Vas1	Cy, Nu
326	Exosome 3'-5' exoribonuclease complex	Csl4 Dis3 Mtr3 Rrp4 Rrp40 Rrp42 Rrp43 Rrp45 Rrp46 Rrp6 Ski6 Ski7	105: Ski2 Ski3 Ski8	Ecm16 Grs1 Lrp1 Rpl4b Ths1 Utp10 Utp20 Yir035c	Cy, No, Nu
327	Complex 327	Sen54 Vac14		Crm1 Rps4b Tao3 Uso1	Va
328	Complex 328	Bud2 Pdr5 Pet9		Cse1 Lys12 Mam33	CP
329	Complex 329	Erg1	111: <i>Erg1 Yhr020w</i>	Dnm1 Gcn20 Pdc1 Sec2 Sec53 Sst2 Tub1	Cy, Nu, Va
330	Complex 330	Fbp26 Ylr345w		Smd1 Snu56 Ssa3 Tao3	Cy
331	Complex 331	Dbp8 Esf1 Hca4 Mvd1 Sik1	69: Ecm16 Krr1 Nop6 72: <i>Hca4</i> Nop58 <i>Sik1</i> 80: Bud21 Imp3	Bfr2 Bms1 Cbf5 Emg1 Ero1 Kre33 Mpp10 Nop1 Pre9 Rok1 Rpl11b Rpl13b Rpl15a Rpl16b Rpl19a Rpl26b Rpl2b Rpl33a Rpl36a Rpl4a Rpl6b Rpp2b Rps14a Rps14b Rps16a Rps18a Rps1b Rps22a Rps24a Rps24b Sof1 Sro9 Utp10 Utp20 Utp30 Utp7	Cy, No
332	Complex 332	Gdh1 Rpl14a		Cdc123 Rpl11b Rpl17b Rpl1a Rpl27a Rpl36a Rpl8b Rpl9a Rps18a Rps20	Cy
333	Septin complex	Bud4 Cdc10 Cdc11 Cdc12 Cdc3 Shs1		Mdh1	Bd, CP
334	Complex 334	Wrs1 Ymr315w		Pdc1 Pep4	Cy, Nu
335	AMP-dependent protein kinase complex	Gal83 Reg1 Sip1 Snf1 Snf4		Ade16 Bmh1 Pak1 Sip2 Ste23	Cy
336	Complex 336	Aha1 Ufd4		Fas2 Lys12 Tdh2 Tif1 Tif2 Tub1 Tub2	Cy, Nu
337	Complex 337	Spt15 Taf1 Taf11 Taf12 Taf3 Taf4 Taf9	146: <i>Taf1 Taf11 Taf12 Taf3 Taf4 Taf9</i>	Med4 Mot1 Myo1 Pdr1 Sec2 Taf5 Taf7 Tao3 Yfl034w	Nu
338	Complex 338	Hua2 Yal048c			
339	Complex 339	Act1 Mgs1			
340	Coatomer COPII complex	Sec13 Sec16 Sec23 Sec24 Sfb2	11: <i>Sec16 Sec23 Sfb2</i> 71: Grh1 <i>Sec24</i>	Cys4 Kar2 Rim1 Rpl10 Rpl7a Sec31 Spt6	ER
341	Complex 341	Nuc1 Ykr079c Ymr099c		Fas2 Ths1	Cy, Nu
342	Cohesin complex	Smc1 Smc3	132: Rnr2 Rnr4 137: <i>Smc1 Smc3</i>	Aro1 Cct8 Eft2 Nop12 Pwp1 Rpl13b Rpl15b Rpl19b Rpl20b Rpl3 Rps6a Vas1 Ypl207w	Cy, Nu
343	Cyclin-dependent protein kinase complex	Cdc28 Cks1 Cln2 Cys4 Sic1		Clb3 Cln1 Met10 Ssc1 Tif1	Cy, Nu
344	Complex 344	Sse1 Tdh2	101: <i>Sse1</i> Tsa1	New1 Pph22 Psa1 Rps0b Rts1 Wrs1 Yal048c	Cy, Nu
345	Complex 345	Aha1 Aro1	34: Trp2 Trp3	Gfa1 Hsc82 Pfk1 Sam1	Cy
346	Complex 346	Sec2		Act1 Gcn5 Med4 Myo1 Ngg1 Pdr1 Sgf29 Spt15 Spt20 Srb2 Taf1 Taf12 Tao3 Yfl034w	Bd, Nu
347	Serine palmitoyltransferase complex	Lcb1 Lcb2	22: <i>Lcb1 Lcb2</i> 94: Kap104 Mir1 108: Sam1 Sam2	Aco1 Bmh2 Gea2 Tub1	ER
348	Complex 348	Ecm5 Iml1 Psd2			
349	Complex 349	Met6 Tkl1		Fip1 Pdc1 Tsa1	Cy, Nu
350	Sec62/Sec63 complex	Sec62 Sec63 Sec66 Sec72		Anp1 Fks1	ER
351	Complex 351	Cdc47 Erg26	90: <i>Cdc47 Erg26</i>		
352	Complex 352	Sam1 Sam2	108: <i>Sam1 Sam2</i>	Aco1 Kap95 Nup2 Rfa3 Rps0b Snf6 Yhr020w	Cy
353	Bro1/Snf7 complex	Bro1 Snf7			Va
354	Complex 354	Dip2 Enp1 Kre33 Krr1 Noc4 Nop14 Rok1 Utp22 Utp6 Utp7 Ygr145w	28: Pwp2 Utp18 Utp21 <i>Utp6</i> 38: Emg1 <i>Noc4 Nop14 Utp22</i> 78: <i>Kre33 Ygr145w</i>	Bfr2 Dbp8 Ero1 Imp3 Mpp10 Nan1 Nop1 Pre9 Rpl15a Rpl19a Rps16a Rps5 Rrp5 Rrp7 Rrp9 Utp30 Utp8 Ylr003c	No

355	Complex 355	Mkt1 Rpl13a Vid24	97: <i>Rpl13a Vid24</i>		Cy
356	Rtg2/Mks1 complex	Mks1 Rtg2		Bmh1 Bmh2 Rvs167	Cy
357	Complex 357	Crm1	94: Kap104 Mir1	Faa4 Gcn1 Hsc82 Lys12 Rna1 Rpt1 Sam1 Tif2 Ura2 Ydj1	Cy, Nu
358	Complex 358	Ssk2	81: Gcn1 Gcn20	Aha1 Cct6 Dnm1 Erg1 Fas2 Grs1 Hsc82 Ilv6 Mkt1 Msn5 Sec53 Ybr025c Yhb1 Ynk1	Cy, Nu
359	Complex 359	Mrs6 Pyk2	104: <i>Mrs6 Pyk2</i>		Cy
360	Complex 360	Tkl1 Tsa1		Gpm1 Met6	Cy
361	Complex 361	Rps0a Rps9b		Eaf5 Mrd1 Rps11a Rps8b	Cy
362	Complex 362	Pmr1 Rvb2 Sec53	100: <i>Rvb2 Sec53</i>	Ils1	Cy
363	Mitochondrial ribosomal large subunit	Img1 Mhr1 Mrp49 Mrpl35 Mrpl8 Mrpl9 Yml6	5: <i>Img1 Mhr1 Mrp49 Mrpl9 Yml6</i> 45: Mrpl10 Mrpl16 Mrpl28 Mrpl7	Aep1 Img2 Mrp20 Mrp7 Mrpl1 Mrpl13 Mrpl15 Mrpl17 Mrpl20 Mrpl22 Mrpl23 Mrpl24 Mrpl27 Mrpl3 Mrpl36 Mrpl4 Mrpl51 Mrpl6 Prp3 Prp31 Shq1 Spp381 Ymr098c Ypl183w-a	Mt
364	SKI complex	Ski2 Ski3 Ski8	105: <i>Ski2 Ski3 Ski8</i>	Asc1 Cdc37 Mtr3 Rpl20a Rpl25 Rpl27a Rpl28 Rpl4b Rpp2b Rps13 Rps22a Rps4a Rps9b Rrp42 Rrp45 Scp160 Skt5	Cy, Nu
365	Rvs161/Rvs167 complex	Rvs161 Rvs167	8: Gyl1 Gyp5 29: <i>Rvs161 Rvs167</i>	Grs1 Idh2 Pfk2 Rpt5 Sec7 Ssa1 Ssb2 Ssk2	Bd, Cs, Cy
366	Complex 366	Ald6 Gsh1 Idh2 New1	89: <i>Ald6 Idh2</i>	Gsp2	Cy
367	Complex 367	Bat1 Bat2	110: Hsp60 Mkt1	Pet56 Ssb2	Cy, Mt
368	Complex 368	Kri1 Rps11b		Asc1 Kre33 Rpl14a Rps10b Rps20 Rps24b Rps4a Rps9b Scp160	Cy, No
369	Complex 369	Fun19 Rps1b			
370	Anaphase-promoting complex, cyclosome	Apc1 Apc2 Apc4 Apc5 Apc9 Cdc16 Cdc23 Cdc27		Yef3	Cy, Nu
371	Complex 371	Nop15	49: Brx1 Erb1 <i>Nop15</i> Nop7 88: Dbp9 Nop4 95: Loc1 Rpl9a	Arx1 Cic1 Dbp10 Drs1 Mak21 Mak5 Mrt4 Nip7 Noc2 Noc3 Nog1 Nsa2 Puf6 Rix1 Rix7 Rlp7 Rpl11b Rpl12a Rpl13a Rpl18b Rpl11b Rpl26b Rpl28 Rpl3 Rpl30 Rpl31a Rpl33b Rpl36b Rpl8b Rpp2b Rps16b Rps18b Rrp12 Rsa3 Spb1 Spb4 Utp20 Yor1 Ytm1	Cy, No, Nu
372	Complex 372	Cin1 Ybl104c		Cdc55 Iml1 Tpd3 Zds1 Zds2	Cy
373	Complex 373	Nfs1 Tdh3 Yer048w-a			Mt
374	COMPASS, histone H3 methyltransferase protein complex	Bre2 Sdc1 Set1 Spp1 Swd1 Swd3		Doa4 Rsm7 Swd2	Nu
375	Complex 375	Pst2 Ybr052c		Fas2 Ssa2 Ynl010w	Cy, Va
376	Complex 376	Mdn1 Nmd5 Tub2		Erg26	Cy, Nu
377	Complex 377	Boi1 Pab1			Cy
378	Complex 378	Arr4 Cdc6			
379	Complex 379	Cym1 Mss51			Mt
380	Retromer subcomplex	Vps29 Vps35 Vps5	20: <i>Vps29 Vps35</i>		Va
381	GPI transamidase complex	Gaa1 Gpi8		Gpi17	ER
382	Complex 382	Stm1 Yil056w			Cy
383	Complex 383	Rpl33b Rpl43a		Dbp10 Dbp2 Imd2 Mak5 Mrps28 Rpl18a Rpl25 Rpl31a Rpl36b Rpl6a Rpp2b Rps18b Sec31 Stm1	Cy
384	Complex 384	Idh2 Lys12	89: <i>Ald6 Idh2</i>	Kap123 Rvb2	Mt
385	Complex 385	Dcs2 Tfb3		Bud20 Gdh1 Rok1 Rpl11b Rpl13a Rpl13b Rpl14b Rpl16a Rpl17b Rpl26b Rpl28 Rpl2a Rpl2b Rpl33b Rpl35b Rpl36a Rpl38 Rpl43a Rpl6a Rpl8a Rps14a Rps16a Rps17b Rps18a Rps1b Rps24a	Cy
386	Complex 386	Rpl1a Tcb3			
387	Complex 387	Ade5,7 Pdr5		Aap1' Clu1 Gea2 Kap104 Tcb3 Usa1	Cy
388	Pab1/Pan2/Pan3 complex	Pab1 Pan2		Pan3	Cy
389	Complex 389	Tdh2 Wrs1	101: Sse1 Tsa1 138: Chc1 Cle1	Acs2 Cmd1 Fba1 Gpm1 New1 Pet9 Pph22 Rpn1 Sec27	Cy, Nu
390	Complex 390	Hir3 Rad53			Nu
391	Complex 391	Aro1 Rad51 Yhb1		Eft1 Met10 Mkt1 Rpa135	Cy
392	Ric1/Rgp1 complex	Rgp1 Ric1	10: <i>Rgp1 Ric1</i>	Tdh3	Go
393	Complex 393	Vip1 Ygr017w		Abf2 Rpl16a Rpl8a Rps0b Rsc4 Rsc58	Cy, Nu
394	Complex 394	Dbp9 Mak21 Mak5 Nop4 Rlp7 Rsa3 Ytm1	88: <i>Dbp9 Nop4</i>	Nop12 Rpl15a Rpl20b Rpl31a Rpl33b Rpl36b Rpl4a Rpp2b Rrp5 Yor1	No
395	Complex 395	Psk1 Taf7			Cy
396	Complex 396	Rpl8a Yer036c	109: <i>Rpl8a Yer036c</i> 113: Fun12 Rpl12b	Rpl3 Rpl6a Rps22a Rps3	Cy

397	Complex 397	Ktr3 Svfl Ypr118w			Cy, Nu
398	Complex 398	Met12 Met13			
399	Complex 399	Ede1 Syp1	<i>127: Ede1 Syp1</i>	Atg26 Lsb3 Sla2 Ydr348c	Bd, CP, Va
400	Heteromeric p24 complex 1	Emp24 Erp1 Erp2 Erv25	25: Erp3 Erp5	Nmd5 Rpl11b Rpt4 Rpt6 Rrp1 Tub3	ER, Va
401	Complex 401	Bud7 Chs5 Chs6 Ymr237w	107: Asc1 Ynl224c	Adh1 Grh1 Rpl1b Rpl5 Skt5 Stm1 Tef1 Vps3 Ykr027w	Go
402	MIND kinetochore complex	Ame1 Dsn1 Mtw1	<i>27: Dsn1 Mtw1</i>	Nkp1 Nsl1	Nu
403	Complex 403	Hir2		Pdr1 Rli1 Rpl12a Rpl1b Rpl33a Rpl35b Rps1b Top2	Cy, Nu
404	Far complex	Far8 Rpl14b		Cdc123 Far3 Tpi1 Ymr134w	Cy, ER
405	Signal recognition particle receptor	Srp101 Srp102	<i>144: Srp101 Srp102</i>		
406	Complex 406	Nop13	35: Cdc33 Tif4632 45: Mrpl10 Mrpl16 Mrpl28 Mrpl7 50: Tps1 Tps2 Tps3	Aep1 Atg26 Brx1 Dbp10 Fas1 Hhf2 Kem1 Kre33 Mrpl15 Mrpl20 Mrpl22 Mrpl23 Mrpl35 Mrpl39 Mrpl51 Mrpl8 Nop12 Nop4 Pol5 Pwp1 Rpl20b Rpl7b Rps14a Rps22b Rps8a Rrp12 Rrp5 Sbp1 Sec1 Shq1 Sro9 Ssb2 Tao3 Tsl1 Ypl183w-a Ypl207w	Cy, Mt, No, Nu
407	TFIIIC complex	Tfc1 Tfc3 Tfc4 Tfc6 Tfc7 Tfc8			Cy, Nu
408	Complex 408	Ist2		Cdc123 Ded1 Glt1 Rpl11a Rpl14b Rpl16a Rpl18a Rpl18b Rpl26b Rpl2b Rpl33a Rpl33b Rpl36b Rps11a Rps16b Rps18a Rps1b Rps6b Rps9a Tif34 Tif35 Tif5	Cy
409	Complex 409	Arx1 Fpr4 Mrt4 Nip7 Nog1 Nog2 Nop2 Nsa2 Nug1 Rpf2 Sda1 Tif6	<i>47: Nog2 Nsa2 Nug1 Tif6</i>		No, Nu
410	Complex 410	Rpl18b Rpl36b		Dps1 Mak5 Nop12 Psa1 Rpl10 Rpl11b Rpl13b Rpl15a Rpl16a Rpl16b Rpl26b Rpl28 Rpl2a Rpl33a Rpl33b Rpl7a Rpl8b Rps14a Rps16b Rps18b Rrp5 Rsa3 Tsa1 Tub3 Vas1 Ytm1	Cy
411	Complex 411	Csn12		Aar2 Brr1 Bud31 Cdc40 Cwc2 Dib1 Ecm2 Hsh155 Hsh49 Isy1 Lea1 Msl1 Prp19 Prp22 Prp4 Prp45 Prp46 Prp9 Rse1 Smd2 Smd3 Snt309 Snu114 Snu23 Ybl104c	Cy, Nu
412	Complex 412	Bud2 Ptm1	2: Dop1 Mon2 92: Cse1 Sec7	Bni1 Cdc48 Crm1 Gcn1 Gea2 Lys12 Mam33 Mir1 Msn5 Pdc1 Pdr5 Pet9 Psa1 Sam1 Ths1 Tub2 Usa1 Ydj1 Yfr044c Yhr020w	Bd, CP, Cy
413	Gim complexes	Gim3 Pac10 Yke2			Cy
414	Complex 414	Pus7 Tan1	<i>112: Pus7 Tan1</i>		Nu
415	Complex 415	Doa4 Dur1,2			
416	Complex 416	Ecm5 Nmd5			Nu
417	Complex 417	Cin1 Rpl43a		Arx1 Dbp10 Fpr4 Nip7 Noc3 Nog1 Nog2 Nop7 Nsa2 Nug1 Rpl11a Rpl25 Rpl30 Rpl31a Rpl36b Rpl38 Sda1 Sxm1	Cy, No, Nu
418	Complex 418	Cwc23 Ylr424w	9: Cwc23 Prp43 Ylr424w 83: Prp39 Smd1 106: Luc7 Smd3 107: Asc1 Ynl224c	Aar2 Brr1 Brr2 Bud31 Cdc33 Cef1 Clf1 Csn12 Cus1 Cwc2 Cwc22 Dib1 Ecm2 Lea1 Lin1 Lsm2 Msl1 Nop1 Prp11 Prp19 Prp21 Prp31 Prp4 Prp45 Prp46 Prp6 Prp8 Prp9 Rps10b Rps11a Rps16b Rps22a Seh1 Smb1 Smd2 Snt309 Snu114 Snu23 Snu71 Sro9 Sto1 Syf1 Ybl104c Yhc1 Ykr022c	Cy, Nu
419	Complex 419	Prs3 Ura7		Por1 Rpl1a Rpl24a Rps5 Ths1 Tom1	Cy
420	DNA polymerase epsilon complex	Dpb2 Dpb4 Pol2	3: Dpb2 Pol2 39: Ctf18 Rfc4 134: Dpb4 Isw2 Itc1	Ded1 Hta2 Imd3 Ioc3 Kap114 Nap1 Pab1 Rfa1 Rfc3 Rps4b Rps6a Rrp8 Top1	Cy, Nu
421	HAP subcomplex of elongator complex	Elp2 Elp3 Elp4 Iki1 Iki3		Hrr25	Cy
422	Complex 422	Mtr10 Nmd5 Usa1		Aap1' Hsc82 Pgi1	Cy, Nu
423	ARP2/3 complex	Arc15 Arc18 Arc19 Arc35 Arc40 Arp2 Arp3		Acc1 Cct5 Cct8 Nog2	Cs
424	Trichostatin A-sensitive class II histone deacetylase complex	Hda1 Hda2 Hda3 Pro2		Msh2	Nu
425	Complex 425	Sbp1	35: Cdc33 Tif4632 115: Sgn1 Ygr250c	Imd4 Kem1 Rps10b Rps20 Rps22b Rps8b Sro9	Cy
426	Complex 426	Rpl30 Ynk1		Rps3	
427	Complex 427	Pep12 Sec17 Vti1 Ykt6	<i>46: Pep12 Ykt6</i>	Akr1 Ste4	

428	RSC, abundant chromatin remodeling complex	Arp7 Npl6 Rsc1 Rsc2 Rsc3 Rsc4 Rsc58 Rsc6 Rsc8 Rsc9 Sfh1 Sth1	24: Arp9 Swi1 Yfl049w 37: Ioc3 Isw1 Mot1 66: Hta2 Htb1 Kap114 136: Yku70 Yku80 147: Snf11 Snf2 Snf5 Snf6 Swi3	Abf2 Hhf1 Hta1 Isw2 Itc1 Pdr1 Reb1 Rsc30 Rtt102 Snf12 Spt15 Stm1 Top1 Vps1 Yil056w	Nu
429	38S HOPS complex	Pep3 Pep5 Vam6 Vps16 Vps33 Vps41 Vps8		Aap1' Prs1	Va
430	Complex 430	Yer007c-a Yjr014w		Por1 Rps1a Rps1b Rps3 Rps4a Rps8a Tfa1	Cy
431	Complex 431	Rli1 Rps7a			
432	Complex 432	Rpl27b Ypl009c		Arx1 Lsg1 Mis1 Nop7 Rpl15a Rpl2b Rpp0 Shm1	Cy, Mt
433	Complex 433	Cys4 Nis1 Ssc1 Yol070c		Nap1	Bd, CP, Cy
434	Complex 434	Arr4 Ded81 Ils1			
435	Complex 435	Chd1 Ssa4		Ckb2 Hot1 Spt16 Ykl088w	Cy, Nu
436	Complex 436	Brr1	106: Luc7 Smd3	Cbc2 Csn12 Cus1 Dib1 Hsh155 Lsm4 Mud1 Prp11 Prp21 Prp38 Prp40 Prp42 Prp6 Prp9 Rse1 Smd1 Smd2 Snp1 Snu56 Snu66 Snu71 Stb1 Tif4631	Cy, Nu
437	Complex 437	Bmh2 Nth1	22: Lcb1 Lcb2	Bmh1 Doa4 Mks1 Psk1 Rtg2 Spt16 Top2	Cy, Nu
438	Complex 438	Lys9 Ynr046w Yol124c		Kar2 Ydr140w Yor164c	Cy, Nu
439	Complex 439	Reb1 Yil056w		Htb1 Rps4b Rsc2 Rsc3 Rsc30 Stm1 Tfc4	Cy, Nu
440	Complex 440	Brx1 Cic1 Mak21 Noc2 Nop4 Spb4 Ytm1	43: <i>Mak21</i> Mak5 Nop2 <i>Ytm1</i> 52: <i>Cic1</i> Ipi1	Drs1 Nip7 Noc3 Nug1 Puf6 Rpl27a Rpl36b	No, Nu
441	Vps34 PtdIns 3 kinase complex II	Vps15 Vps30 Vps34 Vps38			Va
442	Complex 442	Dos2 Kre2			
443	Complex 443	Cct5 Vid27		Acc1 Cct2 Cct6 Cct8 Ilv6 Sit4	Cy
444	Complex 444	Mss51 Ssc1	77: <i>Mss51</i> <i>Ssc1</i> 127: Ede1 Syp1	Cln2 Cym1 Cys4 Eno1 Ist2 Met10 Oye2 Trm3	Cy, Mt
445	Mediator complex, SRB subcomplex of RNA polymerase II	Cse2 Gal11 Med1 Med11 Med6 Med7 Med8 Nut1 Nut2 Pgd1 Rgr1 Sin4 Srb2 Srb4 Srb5 Srb6 Srb7 Srb8 Ssn2	85: <i>Med6</i> Yap1	Ada2 Fab1 Gen5 Hfi1 Med4 Mlp1 Mrpl6 Ngg1 Rpl14a Rpl17b Rpl18a Rpl19b Rpl30 Rpl33a Rpl35b Rpl8a Rps14b Rps17b Sgf29 Sgf73 Snf12 Spt20 Spt3 Spt7 Spt8 Sse2 Ssn3 Ubp8 Vid21	Nu
446	Complex 446	Aro1 Pfk1 Rps0a			Cy
447	Complex 447	Imp3 Ylr003c			
448	Complex 448	Rpl30 Yor1			
449	Complex 449		54: Arp5 Rvb1 116: Hsp82 Lys12 138: Che1 Clc1	Act1 Cct8 Crm1 Cse1 Dps1 Eft2 Fur1 Ies1 Ktr6 Met10 Msn5 Sar1 Ssa4 Tif1 Ufd4 Ura2 Vps13 Ydr341c Ynk1	Cy, Nu
450	Complex 450	Ams1		Dhh1 Rpa135 Rpa190 Rpa49 Rpb10 Rpb5 Rpe31 Rpe40 Rpe82 Rpo31 Sro7	Cy, No, Nu
451	Complex 451	Glt1 Rpl10 Rps5			
452	Complex 452	Mrp8 Pil1 Ylr287c	40: <i>Mrp8</i> <i>Pil1</i>		Cy
453	Complex 453	Gin4 Kcc4	117: <i>Gin4</i> <i>Kcc4</i>	Hmo1 Pgi1 Rps13 Yra1	Bd, Cy
454	Complex 454	Tef4 Ybr025c			Cy
455	Frs1/Frs2 phenylalanyl-tRNA synthetase complex	Frs1 Frs2		Tsc13	
456	Complex 456	Ade5,7 Rps3			
457	Complex 457		87: Nip7 Rpl23a 91: Ebp2 Rpf2	Dbp10 Dbp9 Drs1 Mak21 Nog2 Nop2 Rix7 Rpl11b Rpl1b Rpl25 Rpl8b Rpl9a Rrp1 Spb1 Yer139c	Cy, No, Nu
458	Complex 458	Ilv5 Tdh1		Sar1	
459	RNA polymerase II general transcription factor TFIID	Taf1 Taf11 Taf3 Taf4 Taf5	21: Taf14 Taf2 Taf8 146: <i>Taf1</i> <i>Taf11</i> Taf12 <i>Taf3</i> <i>Taf4</i> Taf9	Hem13 Med4 Myo1 Rps10b Spt15 Spt7 Taf10 Taf6 Taf7 Vps1	Nu
460	Complex 460	Cys3 Rna1			Cy
461	Complex 461	Cct5 Tuf1			
462	Oligosaccharyl-transferase complex	Ost1 Stt3 Swp1 Wbp1		Ero1 Prs1 Tif34	ER
463	Processing peptidase complex	Mas1 Mas2		Dph5 Eft1 Rpl6b	Mt
464	F0/F1 ATP synthase	Atp1 Atp3			Mt
465	Complex 465	Gsh1 Gsp2 New1			Cy
466	Complex 466	Mkt1	110: Hsp60 <i>Mkt1</i>	Gdb1 Gln1 Hsp42 Ilv1 Kap122 Pet56 Por1 Rad51 Rpa135 Rpb2 Vid24 Ydl033c	Cy, Mt, Nu
467	Complex 467	Tbf1 Tor1 Vid22	32: <i>Tbf1</i> <i>Vid22</i> 35: Cdc33 Tif4632 44: Bre5 Ubp3	Nop1 Rrp8 Sik1 Snf2	Cy, Nu
468	Complex 468	Bms1 Bud21 Esf1 Mvd1 Nop6	62: Imp4 Rrp9 Utp13 72: Hca4 Nop58 Sik1 78: Kre33 Ygr145w 80: <i>Bud21</i> Imp3	Bfr2 Cbf5 Dbp8 Ero1 Krr1 Mpp10 Nop14 Rok1 Rpb10 Rpl11b Rpl15a Rpl19a Rpl26b Rpl33a Rpp2b Rps16a Rps22a Sof1 Utp20 Utp30 Utp8	No, Nu

469	Complex 469	Rpl26b Rsa3 Yor1	88: Dbp9 Nop4	Mak21 Mak5 Rpl17b Rpl1b Rpl24a Rpl28 Rpl2a Rpl2b Rpl30 Rpl31a Rpl36b Rpp2b Rps16b Rps18b Rps22b Rps24a Rrp12 Ytm1	Cy, No
470	Ctf18/Rfc2/Rfc3/Rfc4/Rfc5 complex	Elg1 Rfc1 Rfc3 Rfc4 Rfc5	3: Dpb2 Pol2 39: Ctf18 <i>Rfc4</i> 82: Pol1 Pol12 Pri1 Pri2 126: <i>Rfc1 Rfc3 Rfc5</i>	Dpb4 Imd3 Rfc2 Rps0b Rps6b Uso1 Vps13	Cy, Nu
471	Complex 471	Rpl27b Skt5			Cy
472	Complex 472	Dbp10 Dbp9 Nop13 Nop4 Puf6	88: <i>Dbp9 Nop4</i>	Drs1 Nip7 Nug1 Pol5 Rix7 Rlp7 Rpl25 Rpl27a Rpl7b Rrp12	No, Nu
473	Complex 473	Sro7 Ydr314c		Ams1 Ecm29 Rpl6a Rpn1 Rpn10 Rpn11 Rpn12 Rpn3 Rpn6 Rpn8 Rpt2 Ura7	Cy, Nu
474	Complex 474	Hta1 Tos4			
475	Complex 475	Rps25b	78: Kre33 Ygr145w	Hsl1 Mks1 Rpl27a Rpl36a Rps16a Rps17b Rps7a Rps9a Rps9b Rtg2 Sec21 Tif1	Cy
476	Vps27/Hse1 protein complex	Hse1 Vps27			Va
477	Complex 477	Ufd4	103: Pfk1 Pfk2 111: Erg1 Yhr020w	Cct8 Dnm1 Ilv6 Tdh2 Tef1	Cy, Mt, Nu
478	eIF4A	Tif1 Tif2		Gln1 Mcm2 Mks1 Tsc10 Ura7	Cy
479	Complex 479	Gfa1 Msn5 Num1			
480	Complex 480	Sse1 Yhb1		Cys4	Cy
481	Complex 481	Drs2 Pdr5		Tcb3 Usa1	CP
482	Complex 482	Erg1 Tef4	111: <i>Erg1</i> Yhr020w	Pfk1 Sec2 Ybr025c	Cy
483	Complex 483	Dig1 Sap190			
484	Complex 484	Gdi1 Ypt31 Ypt52	15: Skp1 Ymr258c <i>Ypt52</i>	Ypt1	Cy, Nu
485	Complex 485	Bud2 Ccp1 Mrp8	40: <i>Mrp8</i> Pil1		Cy
486	Complex 486	Eft1 Yol098c			Cy
487	Complex 487	Cdc47 Myo5	58: Mlc1 Myo1 68: Cmd1 <i>Myo5</i> 131: Myo4 She2 She3	Act1 Cap1 Cap2 Cdc46 Crn1 Eht1 Mam33 Myo2 Nmd5 Pdr5 Pol5 Rad53	Cs, Nu
488	Complex 488	Ybr025c Ygl039w		Swd3 Ydl033c	Cy
489	Complex 489	Gcn2 Ltv1 Rpm2			Cy
490	Complex 490	Oca1 Siw14 Ycr095c Yhl029c			Cy
491	Complex 491	Shm1 Ypl009c		Ade3 Dbp2 Dcs2 Nog1 Nop7 Rpl11b Rpl13b Rpl16a Rpl16b Rpl18a Rpl18b Rpl25 Rpl26b Rpl27b Rpl2a Rpl2b Rpl33a Rpl33b Rpl35b Rpl36b Rps18a Rps24a Rps3 Tfb3	Cy

* For 13 complexes no core components are defined. Even though the different isoforms share 2/3rds of the components when compared in pairs (i.e. they were merged in a single complex), there is no single protein that is found in at least 2/3rds of all the isoforms.

**In some complexes, modules are found consistently within the various isoforms and are thus fully embedded in cores. In other complexes, only some module components can be found in cores while others are not. This is consequence of the eroding procedure, where constant values are systematically subtracted from socio-affinity scores (see Defining protein complexes from purification data alone in the text for details). For instance, a socio-affinity value of 8 between two proteins in a module may become, for example, 6 during the erosion and thus, the two components might be split. Then, depending on the socio-affinity scores between the other members of the complex, one of the module components might actually be assigned to more isoforms than the other and thus, finally, it will be considered a component of the complex core.

The module components that overlap with cores are indicated in italics.