

Size	Complex name	MCL	MCODE	DPCLUS	CMC	COACH	RRW	NWE	PPSampler1	PPSampler2
2	6-phosphofructokinase complex	0	0	0	0	0.45	0	0.82	0	1
	Ribosome-associated complex (RAC)	0.50	0	0	0.47	0	0	0.82	0	1
	Fig4p/Vac14p complex	0.54	0	0	0.53	0.71	0	0.82	0	1
	anthranilate synthase complex	0.71	0	0	0	0	0	0.82	0	1
3	Rad17p/Ddc1p/Mec3p complex	0.71	0	0	0	0.52	0.82	0.67	0.77	1
	chromatin silencing complex	0.61	0	0	0	0.61	0	0	0.65	1
	karyopherin docking subcomplex of the Nuclear Pore Complex (NPC)	0.61	0	0.65	0.61	0	0	0	0	1
	DNA polymerase delta complex	0.87	0	0.87	0	0.47	0.82	0.87	0.82	1
	NatA complex	0.65	0	0	0	0.61	0.82	0.87	0	1
	Kel1p/Kel2p/Lte1p complex	0.77	0	0	0	0	0.82	0	0.71	1
	Apg12p/Apg5p/Apg16 multimeric complex	0.50	0	0.52	0	0.55	0.82	0.87	0.87	1
	Polzeta-Rev1p complex	0.71	0	0	0	0	0.82	0.67	0	1
	methionyl glutamyl tRNA synthetase complex	0.71	0	0	0	0	0	0.87	0.50	1
	RNA Degradation Complex	0.61	0	0	0.58	0	0.82	0.67	0	1
	Atg20p/Snx4p/Snx41p complex	0.65	0	0.71	0	0	0.82	0.67	0	1
	RNA polymerase I transcription factor complex	0.55	0.61	0.77	0	0.87	0.82	0.87	0	1
4	alpha DNA polymerase:primase complex	0.75	0	0.52	0	0.55	0.71	0.75	0.75	1
	alpha,alpha-trehalose-phosphate synthase complex (UDP-forming)	0.50	0	0.50	0	0.53	0	0.89	0.58	1
	ARGR complex	0.55	0	0.82	0	0.63	0.71	0.75	0.87	1
	Sec62p/Sec63p complex	0.76	0	0.82	0	0.82	0.87	0.75	0.82	1
	Ndc80p complex	0.67	0	0.76	0	0.76	0.87	0	0.82	1
	Nsp1p complex	0.55	0	0.55	0	0.63	0.87	0	0	1
	AP-3 adaptor complex	0.71	0.87	0.71	0	0.82	0.71	0.58	0.71	1
5	eEF1	0.67	0	0.68	0	0.73	0.63	0.52	0.52	1
	AP-1 adaptor complex	0.51	0	0.71	0	0.71	0.77	0	0	1
	alpha-1,6-mannosyltransferase complex (Anp1p/Mnn9p)	0.53	0	0.71	0	0.75	0.77	0	0.62	1
6	Nup84p complex	0.52	0	0.71	0	0.72	0.91	0	0.46	1
	Signal recognition particle (SRP)	0.82	0.77	0.82	0	0.77	0.82	0.93	0.93	1
	prefoldin complex	0.65	0	0.87	0	0.77	0	0.47	0.73	1
7	Arp2/3 protein complex	0.61	0	0.68	0	0.61	0.65	0	0.61	1
	Set3p complex	0.71	0	0.73	0	0.84	0.53	0.86	0.94	1
	nucleotide-excision repair factor 3 complex	0.54	0	0.84	0	0.71	0.53	0.57	0.57	1
9	oligosaccharyl transferase complex	0.77	0.52	0.80	0	0.75	0.88	0	0.82	1
14	20S proteasome	0.78	0	0.84	0	0.91	0.53	0	0.69	1
25	Kornberg's mediator (SRB) complex	0.88	0.50	0.93	0	0.93	0.53	0	0.83	1