

Complex	Median $k_{deg}(h^{-1})$	Variance	<i>P</i>	# Members
20S core proteasome ⁵⁻⁷	0.013	0.00037	0.000055	15
17S U2 snRNP/SF3b (spliceosome) ⁸	0.015	0.00369	0.00073	22
Macromolecular tRNA synthetase complex ⁹	0.013	0.00042	0.002559	10
chaperonin-containing t-complex ¹⁰	0.014	0.00044	0.004181	9
Mammalian translation initiation factor 3 (eIF3) ¹¹	0.019	0.00327	0.007149	11
COP9 signalosome complex ^{12, 13}	0.013	0.00114	0.011092	8
19S proteasome regulatory cap ⁵⁻⁷	0.014	0.00203	0.01126	10
COPI coatomer complex ¹⁴	0.016	0.00056	0.011763	7
ER complex that binds nascent proteins ¹⁵	0.015	0.00034	0.012521	7
G-protein complex ¹⁶	0.024	0.00159	0.017752	8
Aminoacyl-tRNA synthetase : EF-1 complex ¹⁷	0.012	0.00011	0.019523	4
Messenger ribonucleoprotein complex ¹⁸	0.016	0.00551	0.020048	17
Histone deacetylase / nucleosome remodeling activities complex ¹⁹	0.024	0.00689	0.022413	9
Histone deacetylase-containing corepressor complexes ²⁰	0.025	0.00587	0.027024	9
DNA-dependent protein kinase complex ²¹	0.014	0.00063	0.029323	5
Double-stranded RNA activated protein kinase PKR complex ²²	0.013	0.0014	0.03585	5
Nonsense-Mediated mRNA Decay complex ²³	0.020	0.00187	0.04318	7
Secretory glycoprotein associated ER chaperone complex ²⁴	0.015	0.00025	0.044901	4
Asialoglycoprotein receptor bound HSP complex ²⁵	0.014	0.00064	0.045527	4