

GOTERM category	Term	PValue	Fold Enrichment	Bonferroni	Benjamini	FDR	
Biological Process	GO:0006414~translational elongation	4,13E-131	81,28	2,67E-128	2,67E-128	6,16E-128	
	GO:0006412~translation	2,87E-90	25,85	1,85E-87	9,26E-88	4,28E-87	
	GO:0042274~ribosomal small subunit biogenesis	9,90E-18	105,11	6,39E-15	2,13E-15	1,48E-14	
	GO:0042254~ribosome biogenesis	1,11E-17	18,01	7,15E-15	1,79E-15	1,65E-14	
	GO:0006396~RNA processing	1,07E-14	6,13	6,89E-12	1,38E-12	1,59E-11	
	GO:0022613~ribonucleoprotein complex biogenesis	1,31E-14	12,20	8,46E-12	1,41E-12	1,95E-11	
	GO:0006364~rRNA processing	3,74E-14	18,85	2,42E-11	3,45E-12	5,58E-11	
	GO:0016072~rRNA metabolic process	6,92E-14	18,07	4,47E-11	5,59E-12	1,03E-10	
	GO:0042273~ribosomal large subunit biogenesis	3,33E-13	92,50	2,15E-10	2,39E-11	4,98E-10	
	GO:0034470~ncRNA processing	5,95E-11	9,89	3,85E-08	3,85E-09	8,88E-08	
	GO:0034660~ncRNA metabolic process	1,09E-09	8,04	7,06E-07	6,42E-08	1,63E-06	
	GO:0043488~regulation of mRNA stability	3,33E-05	26,28	2,13E-02	1,79E-03	4,97E-02	
	GO:0045103~intermediate filament-based process	3,33E-05	26,28	2,13E-02	1,79E-03	4,97E-02	
	GO:0006413~translational initiation	3,95E-05	15,42	2,52E-02	1,96E-03	5,89E-02	
	GO:0043487~regulation of RNA stability	4,78E-05	24,09	3,04E-02	2,20E-03	7,12E-02	
	GO:0042255~ribosome assembly	7,06E-05	46,25	4,46E-02	3,03E-03	1,05E-01	
	GO:0016071~mRNA metabolic process	7,81E-05	4,06	4,92E-02	3,15E-03	1,16E-01	
	GO:0030705~cytoskeleton-dependent intracellular transport	8,01E-05	13,34	5,04E-02	3,04E-03	1,19E-01	
	GO:0010608~posttranscriptional regulation of gene expression	8,30E-05	5,48	5,22E-02	2,98E-03	1,24E-01	
	GO:0042257~ribosomal subunit assembly	2,17E-04	115,62	1,31E-01	7,37E-03	3,24E-01	
	GO:0043489~RNA stabilization	2,59E-04	30,83	1,54E-01	8,34E-03	3,86E-01	
	GO:0048255~mRNA stabilization	2,59E-04	30,83	1,54E-01	8,34E-03	3,86E-01	
	GO:0045104~intermediate filament cytoskeleton organization	6,30E-04	23,12	3,34E-01	1,92E-02	9,35E-01	
	GO:0033119~negative regulation of RNA splicing	7,17E-04	69,37	3,71E-01	2,08E-02	1,06E+00	
	GO:0006397~mRNA processing	1,76E-03	3,60	6,79E-01	4,83E-02	2,59E+00	
	Cellular Component	GO:0022626~cytosolic ribosome	5,46E-117	82,85	9,17E-115	9,17E-115	6,64E-114
		GO:0005840~ribosome	4,72E-103	36,66	7,93E-101	3,97E-101	5,75E-100
		GO:0033279~ribosomal subunit	1,40E-102	54,09	2,36E-100	7,86E-101	1,71E-99
		GO:0030529~ribonucleoprotein complex	1,40E-100	18,61	2,36E-98	5,89E-99	1,71E-97
		GO:0044445~cytosolic part	4,75E-94	44,85	7,98E-92	1,60E-92	5,78E-91
		GO:0022627~cytosolic small ribosomal subunit	3,97E-87	85,21	6,67E-85	1,11E-85	4,84E-84
		GO:0005829~cytosol	1,11E-85	6,81	1,86E-83	2,66E-84	1,35E-82
		GO:0022625~cytosolic large ribosomal subunit	4,80E-83	84,09	8,06E-81	1,01E-81	5,84E-80
GO:0043232~intracellular non-membrane-bounded organelle		4,43E-82	4,19	7,43E-80	8,26E-81	5,39E-79	
GO:0043228~non-membrane-bounded organelle		4,43E-82	4,19	7,43E-80	8,26E-81	5,39E-79	
GO:0015935~small ribosomal subunit		7,90E-80	55,79	1,33E-77	1,33E-78	9,62E-77	
GO:0015934~large ribosomal subunit		1,13E-48	52,46	1,89E-46	1,72E-47	1,37E-45	
GO:0005730~nucleolus		4,45E-09	3,97	7,47E-07	6,23E-08	5,42E-06	
GO:0031981~nuclear lumen		1,74E-05	2,28	2,93E-03	2,25E-04	2,12E-02	
GO:0060053~neurofilament cytoskeleton		9,00E-05	42,61	1,50E-02	1,08E-03	1,10E-01	
GO:0070013~intracellular organelle lumen		1,36E-04	1,98	2,26E-02	1,52E-03	1,65E-01	
GO:0043233~organelle lumen		2,08E-04	1,93	3,44E-02	2,19E-03	2,53E-01	
GO:0031974~membrane-enclosed lumen		2,99E-04	1,89	4,90E-02	2,95E-03	3,64E-01	
GO:0005844~polysome		5,80E-04	23,67	9,28E-02	5,40E-03	7,04E-01	
GO:0005883~neurofilament		1,26E-03	53,26	1,91E-01	1,11E-02	1,52E+00	
GO:0005681~spliceosome		1,45E-03	5,65	2,16E-01	1,21E-02	1,75E+00	
Molecular Function		GO:0003735~structural constituent of ribosome	1,58E-100	43,00	2,36E-98	2,36E-98	1,89E-97
		GO:0005198~structural molecule activity	1,50E-73	13,21	2,23E-71	1,11E-71	1,78E-70
	GO:0003723~RNA binding	4,13E-66	11,37	6,15E-64	2,05E-64	4,93E-63	
	GO:0003729~mRNA binding	8,66E-19	26,97	1,29E-16	3,23E-17	1,03E-15	
	GO:0019843~rRNA binding	3,90E-12	36,10	5,82E-10	1,16E-10	4,66E-09	
	GO:0003725~double-stranded RNA binding	2,54E-04	15,86	3,71E-02	6,28E-03	3,02E-01	
	GO:0005200~structural constituent of cytoskeleton	6,72E-04	8,49	9,54E-02	1,42E-02	7,99E-01	
	GO:0008143~poly(A) RNA binding	2,40E-03	39,26	3,01E-01	4,38E-02	2,83E+00	
	GO:0045182~translation regulator activity	2,53E-03	14,44	3,15E-01	4,11E-02	2,98E+00	
	GO:0070717~poly-purine tract binding	3,07E-03	34,90	3,67E-01	4,48E-02	3,60E+00	