

GO ID	GO Term	Ontology	Number Observed	Total Number	P-value
GO:0005830	Cytosolic ribosome (sensu Eukaryota)	C	25	39	1.07e-12
GO:0030529	Ribonucleoprotein complex	C	48	126	6.12e-09
GO:0006950	Response to stress	P	72	229	1.61e-08
GO:0044445	Cytosolic part	C	27	55	1.61e-08
GO:0003735	Structural constituent of ribosome	F	29	65	1.4e-07
GO:0005829	Cytosol	C	47	132	1.4e-07
GO:0005840	Ribosome	C	30	69	1.63e-07
GO:0044421	Extracellular region part	C	58	180	2.48e-07
GO:0003723	RNA binding	F	56	175	6.61e-07
GO:0005737	Cytoplasm	C	217	1014	1.1e-06
GO:0005615	Extraceullar space	C	41	118	3.63e-06
GO:0006412	Translation	P	46	140	5.28e-06
GO:0051082	Unfolded protein binding	F	18	36	9.69e-06
GO:0043228	Non-membrane-bound organelle	C	93	370	3.42e-05
GO:0043232	Intracellular non-membrane-bound organelle	C	93	370	3.42e-05
GO:0016282	Eukaryotic 43S preinitiation complex	C	14	22	7.22e-05
GO:0005843	Cytosolic small ribosomal subunit (sensu Eukaryota)	C	12	17	8.38e-05
GO:0016283	Eukaryotic 48S initiation complex	C	12	17	8.38e-05
GO:0005842	Cytosolic large ribosomal subunit (sensu Eukaryota)	C	13	21	0.000253
GO:0008283	Cell proliferation	P	54	194	0.000372
GO:0015935	Small ribosomal subunit	C	12	19	0.000431
GO:0009611	Response to wounding	P	31	93	0.000513
GO:0032991	Macromolecular complex	C	110	482	0.00057
GO:0000074	Regulation of progression through cell cycle	P	46	161	0.000784