



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0005739	mitochondrion	1556	4.2e-135
2	REAC	REAC:R-HSA-72766	Translation	281	2.0e-131
3	GO:CC	GO:0044391	ribosomal subunit	183	1.4e-117
4	GO:CC	GO:0005740	mitochondrial envelope	740	1.9e-111
5	GO:BP	GO:1901566	organonitrogen compound biosynthetic process	1653	7.5e-106
6	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	2.1e-105
7	GO:BP	GO:0006412	translation	685	1.5e-95
8	GO:BP	GO:0043043	peptide biosynthetic process	711	1.6e-94
9	REAC	REAC:R-HSA-1430728	Metabolism	1832	7.0e-90
0	KEGG	KEGG:03010	Ribosome	128	1.3e-86
1	KEGG	KEGG:05012	Parkinson disease	234	4.6e-61
12	KEGG	KEGG:05016	Huntington disease	274	1.7e-57
13	GO:MF	GO:0003735	structural constituent of ribosome	191	1.1e-46
4	GO:MF	GO:0015453	oxidoreduction-driven active transmembrane transporter activity	55	3.2e-31
5	GO:MF	GO:0003723	RNA binding	1885	4.9e-31

g:Profiler (biit.cs.ut.ee/gprofiler)