



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0005739	mitochondrion	1556	7.1e-140
2	REAC	REAC:R-HSA-72766	Translation	281	4.4e-129
3	GO:BP	GO:1901566	organonitrogen compound biosynthetic process	1653	6.2e-116
4	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	9.0e-113
5	GO:CC	GO:0005740	mitochondrial envelope	740	8.2e-112
6	GO:CC	GO:0031975	envelope	1164	7.7e-111
7	REAC	REAC:R-HSA-8953854	Metabolism of RNA	626	4.1e-104
8	GO:BP	GO:0006412	translation	685	5.2e-98
9	GO:BP	GO:0043043	peptide biosynthetic process	711	1.0e-95
0	KEGG	KEGG:03010	Ribosome	128	1.1e-77
1	KEGG	KEGG:05012	Parkinson disease	234	4.0e-58
12	KEGG	KEGG:05016	Huntington disease	274	3.4e-55
13	GO:MF	GO:0003723	RNA binding	1885	5.1e-43
14	GO:MF	GO:0003735	structural constituent of ribosome	191	1.3e-39
15	GO:MF	GO:0015453	oxidoreduction-driven active transmembrane transporter activity	55	3.0e-29

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