



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0098588	bounding membrane of organelle	1933	3.3e-54
2	GO:CC	GO:0005739	mitochondrion	1556	2.9e-48
3	REAC	REAC:R-HSA-1430728	Metabolism	1832	6.3e-45
4	GO:CC	GO:0030054	cell junction	1916	3.7e-44
5	GO:BP	GO:0044281	small molecule metabolic process	1627	2.5e-40
6	GO:MF	GO:0019899	enzyme binding	1888	1.4e-39
7	GO:BP	GO:1901566	organonitrogen compound biosynthetic process	1653	7.6e-38
8	GO:BP	GO:0044248	cellular catabolic process	1938	7.7e-36
9	KEGG	KEGG:01100	Metabolic pathways	1334	2.3e-35
10	GO:MF	GO:1901265	nucleoside phosphate binding	1974	2.1e-33
11	GO:MF	GO:0000166	nucleotide binding	1973	3.6e-33
12	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	1.6e-25
13	REAC	REAC:R-HSA-9675108	Nervous system development	531	2.7e-19
14	KEGG	KEGG:04144	Endocytosis	238	2.6e-10
15	KEGG	KEGG:04714	Thermogenesis	207	3.5e-10

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