



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0005739	mitochondrion	1556	8.5e-54
2	GO:BP	GO:0044248	cellular catabolic process	1938	1.9e-49
3	GO:CC	GO:0031967	organelle envelope	1164	8.7e-43
4	GO:CC	GO:0031975	envelope	1164	8.7e-43
5	GO:BP	GO:1901575	organic substance catabolic process	1896	2.4e-42
6	GO:BP	GO:0046907	intracellular transport	1572	2.0e-38
7	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	1.3e-37
8	GO:MF	GO:0019899	enzyme binding	1888	1.2e-31
9	GO:MF	GO:0000166	nucleotide binding	1973	4.6e-30
10	GO:MF	GO:1901265	nucleoside phosphate binding	1974	5.0e-30
11	REAC	REAC:R-HSA-1430728	Metabolism	1832	3.7e-29
12	REAC	REAC:R-HSA-597592	Post-translational protein modification	1257	4.1e-29
13	KEGG	KEGG:01100	Metabolic pathways	1334	4.0e-22
14	KEGG	KEGG:05022	Pathways of neurodegeneration – multiple diseases	421	1.1e-16
15	KEGG	KEGG:05016	Huntington disease	274	4.5e-15

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