



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-8953854	Metabolism of RNA	626	7.2e-11
2	GO:BP	GO:0031324	negative regulation of cellular metabolic process	1963	2.8e-10
3	GO:CC	GO:0016604	nuclear body	796	8.4e-10
4	GO:MF	GO:0019899	enzyme binding	1888	3.4e-09
5	GO:BP	GO:0050684	regulation of mRNA processing	130	7.0e-08
6	REAC	REAC:R-HSA-1643685	Disease	1562	1.1e-07
7	GO:BP	GO:1903311	regulation of mRNA metabolic process	280	1.1e-07
8	REAC	REAC:R-HSA-422475	Axon guidance	507	1.3e-07
9	GO:CC	GO:1902494	catalytic complex	1628	5.3e-07
10	GO:MF	GO:0000166	nucleotide binding	1973	6.9e-07
11	GO:MF	GO:1901265	nucleoside phosphate binding	1974	7.0e-07
12	GO:CC	GO:0030054	cell junction	1916	1.4e-06
13	KEGG	KEGG:03010	Ribosome	128	6.2e-04
14	KEGG	KEGG:05171	Coronavirus disease – COVID–19	184	6.1e-03
15	KEGG	KEGG:03040	Spliceosome	133	6.5e-03

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