



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0006259	DNA metabolic process	973	2.3e-09
2	GO:CC	GO:0016604	nuclear body	796	4.6e-09
3	GO:BP	GO:0006403	RNA localization	187	5.0e-09
4	GO:CC	GO:1902494	catalytic complex	1628	5.3e-09
5	GO:CC	GO:0016607	nuclear speck	394	8.8e-08
6	GO:BP	GO:0033043	regulation of organelle organization	1106	8.9e-08
7	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	1.3e-07
8	GO:MF	GO:0003723	RNA binding	1885	1.4e-06
9	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	2.6e-06
10	REAC	REAC:R-HSA-5617833	Cilium Assembly	188	6.6e-06
11	GO:MF	GO:0019899	enzyme binding	1888	7.8e-05
12	GO:MF	GO:0030554	adenyl nucleotide binding	1523	1.7e-04
13	KEGG	KEGG:03013	Nucleocytoplasmic transport	98	2.7e-03
14	KEGG	KEGG:00640	Propanoate metabolism	30	1.7e-02
15	KEGG	KEGG:00620	Pyruvate metabolism	39	1.8e-02

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