



id	source	term_id	term_name	term_size	p_value
1	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	7.6e-49
2	GO:MF	GO:0003690	double-stranded DNA binding	1415	2.5e-46
3	GO:MF	GO:0043565	sequence-specific DNA binding	1419	8.7e-46
4	GO:MF	GO:1990837	sequence-specific double-stranded DNA binding	1321	1.7e-43
5	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	2.9e-40
6	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	8.0e-36
7	REAC	REAC:R-HSA-212436	Generic Transcription Pathway	1130	3.8e-30
8	GO:CC	GO:0016604	nuclear body	796	1.4e-22
9	GO:BP	GO:0045935	positive regulation of nucleobase-containing compound metabolic process	1798	3.4e-20
10	GO:CC	GO:0005815	microtubule organizing center	787	8.8e-20
11	GO:CC	GO:0005813	centrosome	596	1.0e-18
12	GO:BP	GO:0051254	positive regulation of RNA metabolic process	1610	5.2e-18
13	GO:BP	GO:0007049	cell cycle	1655	1.6e-16
14	KEGG	KEGG:03040	Spliceosome	133	1.5e-05
15	KEGG	KEGG:03013	Nucleocytoplasmic transport	98	1.3e-04

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