



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	2.6e-47
2	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	6.9e-38
3	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	3.1e-36
4	GO:MF	GO:0003690	double-stranded DNA binding	1415	2.8e-33
5	GO:MF	GO:0043565	sequence-specific DNA binding	1419	3.2e-31
6	GO:MF	GO:0000976	transcription cis-regulatory region binding	1272	1.1e-30
7	REAC	REAC:R-HSA-212436	Generic Transcription Pathway	1130	1.1e-28
8	GO:CC	GO:0016604	nuclear body	796	6.7e-24
9	GO:CC	GO:0016607	nuclear speck	394	6.7e-17
10	GO:BP	GO:0045934	negative regulation of nucleobase-containing compound metabolic process	1332	4.2e-13
11	GO:BP	GO:0051253	negative regulation of RNA metabolic process	1219	2.7e-12
12	GO:BP	GO:0031327	negative regulation of cellular biosynthetic process	1375	3.9e-12
13	GO:CC	GO:1990234	transferase complex	798	1.3e-10
14	KEGG	KEGG:03013	Nucleocytoplasmic transport	98	1.3e-05
15	KEGG	KEGG:03040	Spliceosome	133	3.2e-04

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