



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	4.8e-28
2	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	2.5e-22
3	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	2.9e-20
4	GO:MF	GO:0003690	double-stranded DNA binding	1415	4.5e-20
5	GO:MF	GO:0043565	sequence-specific DNA binding	1419	7.7e-19
6	GO:MF	GO:0001067	transcription regulatory region nucleic acid binding	1274	3.7e-18
7	REAC	REAC:R-HSA-212436	Generic Transcription Pathway	1130	3.5e-17
8	GO:CC	GO:0016604	nuclear body	796	7.7e-17
9	GO:CC	GO:0016607	nuclear speck	394	7.5e-13
10	GO:BP	GO:0045934	negative regulation of nucleobase-containing compound metabolic process	1332	1.1e-11
11	GO:BP	GO:0031324	negative regulation of cellular metabolic process	1963	1.8e-11
12	GO:BP	GO:0010558	negative regulation of macromolecule biosynthetic process	1324	1.3e-09
13	GO:CC	GO:1902494	catalytic complex	1628	3.6e-07
14	KEGG	KEGG:03250	Viral life cycle – HIV–1	60	2.3e-05
15	KEGG	KEGG:03013	Nucleocytoplasmic transport	98	5.5e-04