



id	source	term_id	term_name	term_size	p_value
1	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	3.5e-37
2	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	1.8e-25
3	GO:MF	GO:0000976	transcription cis-regulatory region binding	1272	1.9e-23
4	GO:MF	GO:0001067	transcription regulatory region nucleic acid binding	1274	2.1e-23
5	GO:MF	GO:0000977	RNA polymerase II transcription regulatory region sequence-specific DNA binding	1186	2.2e-23
6	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	2.0e-22
7	REAC	REAC:R-HSA-212436	Generic Transcription Pathway	1130	3.6e-22
8	GO:CC	GO:1902494	catalytic complex	1628	8.9e-21
9	GO:CC	GO:1990234	transferase complex	798	2.7e-17
10	GO:CC	GO:0140535	intracellular protein-containing complex	869	4.0e-17
11	GO:BP	GO:0044265	cellular macromolecule catabolic process	928	3.2e-16
12	GO:BP	GO:0006511	ubiquitin-dependent protein catabolic process	623	5.1e-16
13	GO:BP	GO:0019941	modification-dependent protein catabolic process	633	9.2e-16
14	KEGG	KEGG:04120	Ubiquitin mediated proteolysis	129	7.4e-06
15	KEGG	KEGG:04137	Mitophagy – animal	68	7.6e-05