



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-8953854	Metabolism of RNA	626	3.3e-57
2	GO:CC	GO:0005694	chromosome	1638	4.1e-43
3	GO:BP	GO:0006974	cellular response to DNA damage stimulus	827	2.0e-39
4	REAC	REAC:R-HSA-1640170	Cell Cycle	640	2.9e-39
5	GO:BP	GO:0006259	DNA metabolic process	973	2.9e-38
6	GO:BP	GO:0007049	cell cycle	1655	1.2e-37
7	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	1.7e-37
8	GO:CC	GO:0098687	chromosomal region	366	1.8e-30
9	GO:MF	GO:0140640	catalytic activity, acting on a nucleic acid	574	5.1e-29
10	GO:CC	GO:1902494	catalytic complex	1628	1.8e-27
11	GO:MF	GO:0003723	RNA binding	1885	6.9e-25
12	GO:MF	GO:0003690	double-stranded DNA binding	1415	1.1e-17
13	KEGG	KEGG:03040	Spliceosome	133	1.2e-13
14	KEGG	KEGG:03030	DNA replication	34	1.4e-13
15	KEGG	KEGG:03013	Nucleocytoplasmic transport	98	2.0e-12