



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0003712	transcription coregulator activity	527	1.2e-06
2	GO:MF	GO:0003682	chromatin binding	603	1.4e-04
3	GO:MF	GO:0140297	DNA-binding transcription factor binding	490	1.2e-03
4	GO:MF	GO:0016922	nuclear receptor binding	138	1.9e-04
5	GO:CC	GO:0031981	nuclear lumen	5524	4.7e-17
6	GO:CC	GO:0016514	SWI/SNF complex	30	1.4e-04
7	GO:CC	GO:0000785	chromatin	1371	8.8e-06
9	GO:BP	GO:0006357	regulation of transcription by RNA polymerase II	2572	2.1e-04
10	GO:BP	GO:0006325	chromatin organization	843	1.2e-05
12	KEGG	KEGG:03082	ATP-dependent chromatin remodeling	116	5.5e-03
13	REAC	REAC:R-HSA-4839726	Chromatin organization	271	1.4e-02
14	REAC	REAC:R-HSA-9006931	Signaling by Nuclear Receptors	290	4.9e-02