



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0003712	transcription coregulator activity	527	4.2e-20
2	GO:MF	GO:0003682	chromatin binding	603	1.4e-16
3	GO:MF	GO:0140297	DNA-binding transcription factor binding	490	1.3e-12
4	GO:MF	GO:0016922	nuclear receptor binding	138	1.8e-11
5	GO:CC	GO:0031981	nuclear lumen	5524	1.0e-16
6	GO:CC	GO:0016514	SWI/SNF complex	30	1.0e-14
7	GO:CC	GO:0000785	chromatin	1371	6.6e-13
8	GO:BP	GO:0141193	nuclear receptor-mediated signaling pathway	179	1.4e-12
9	GO:BP	GO:0006357	regulation of transcription by RNA polymerase II	2572	7.6e-12
10	GO:BP	GO:0006325	chromatin organization	843	1.3e-10
11	GO:BP	GO:0030518	nuclear receptor-mediated steroid hormone signaling pathway	123	6.8e-10
12	KEGG	KEGG:03082	ATP-dependent chromatin remodeling	116	9.6e-08
13	REAC	REAC:R-HSA-4839726	Chromatin organization	271	3.9e-08
14	REAC	REAC:R-HSA-9006931	Signaling by Nuclear Receptors	290	6.4e-05