



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0048870	cell motility	1470	1.4e-13
2	GO:BP	GO:0022008	neurogenesis	1548	2.4e-13
3	GO:BP	GO:0048646	anatomical structure formation involved in morphogenesis	1021	3.3e-12
4	GO:CC	GO:0005929	cilium	635	4.2e-10
5	GO:CC	GO:0031514	motile cilium	201	4.7e-08
6	GO:CC	GO:0005912	adherens junction	168	1.5e-07
7	REAC	REAC:R-HSA-5083635	Defective B3GALTL causes PpS	34	2.0e-06
8	REAC	REAC:R-HSA-5173214	O-glycosylation of TSR domain-containing proteins	35	2.6e-06
9	REAC	REAC:R-HSA-9012999	RHO GTPase cycle	419	4.5e-06
10	GO:MF	GO:1990837	sequence-specific double-stranded DNA binding	1321	6.1e-06
11	GO:MF	GO:0008237	metallopeptidase activity	149	8.5e-06
12	GO:MF	GO:0000977	RNA polymerase II transcription regulatory region sequence–specific DNA binding	1186	9.0e-06
13	KEGG	KEGG:05165	Human papillomavirus infection	294	7.9e-05
14	KEGG	KEGG:04330	Notch signaling pathway	55	2.2e-04
15	KEGG	KEGG:05224	Breast cancer	132	4.5e-04