



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0009887	animal organ morphogenesis	907	4.9e-21
2	GO:MF	GO:1990837	sequence-specific double-stranded DNA binding	1321	6.0e-18
3	GO:MF	GO:0000981	DNA-binding transcription factor activity, RNA polymerase II-specific	1131	1.2e-17
4	GO:MF	GO:0000977	RNA polymerase II transcription regulatory region sequence–specific DNA binding	1186	6.2e-17
5	GO:BP	GO:0009888	tissue development	1647	4.6e-16
6	GO:BP	GO:0035295	tube development	957	4.6e-15
7	GO:CC	GO:0000785	chromatin	1112	6.2e-12
8	GO:CC	GO:0005929	cilium	635	1.2e-07
9	GO:CC	GO:0005694	chromosome	1638	2.1e-07
10	REAC	REAC:R-HSA-5083635	Defective B3GALTL causes PpS	34	1.1e-05
11	REAC	REAC:R-HSA-5173214	O-glycosylation of TSR domain-containing proteins	35	1.3e-05
12	KEGG	KEGG:04550	Signaling pathways regulating pluripotency of stem cells	129	1.3e-04
13	REAC	REAC:R-HSA-3906995	Diseases associated with O–glycosylation of proteins	53	2.8e-04
14	KEGG	KEGG:04310	Wnt signaling pathway	150	9.1e-04
15	KEGG	KEGG:04015	Rap1 signaling pathway	199	6.4e-03