



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0000166	nucleotide binding	1973	2.1e-20
2	GO:MF	GO:1901265	nucleoside phosphate binding	1974	2.2e-20
3	GO:MF	GO:0017076	purine nucleotide binding	1835	1.0e-19
4	GO:BP	GO:0016310	phosphorylation	1653	7.7e-19
5	GO:BP	GO:0018193	peptidyl-amino acid modification	1120	1.1e-18
6	GO:BP	GO:0006468	protein phosphorylation	1379	1.4e-17
7	GO:CC	GO:0030054	cell junction	1916	1.7e-17
8	GO:CC	GO:0045202	synapse	1219	3.2e-15
9	GO:CC	GO:0043005	neuron projection	1254	4.5e-13
10	REAC	REAC:R-HSA-1643685	Disease	1562	5.7e-10
11	REAC	REAC:R-HSA-9716542	Signaling by Rho GTPases, Miro GTPases and RHOBTB3	662	2.9e-08
12	REAC	REAC:R-HSA-194315	Signaling by Rho GTPases	649	3.1e-08
13	KEGG	KEGG:04727	GABAergic synapse	78	1.4e-06
14	KEGG	KEGG:04724	Glutamatergic synapse	107	5.0e-06
15	KEGG	KEGG:05169	Epstein-Barr virus infection	170	4.8e-05