



id	source	term_id	term name	term_size	p value
1	GO:MF	GO:0030554	adenyl nucleotide binding	1523	2.2e-14
2	GO:MF	GO:0000166	nucleotide binding	1973	3.2e-14
3	GO:MF	GO:1901265	nucleoside phosphate binding	1974	3.3e-14
4	GO:BP	GO:0044281	small molecule metabolic process	1627	1.9e-11
5	GO:BP	GO:0016570	histone modification	434	2.5e-10
6	GO:BP	GO:0018193	peptidyl-amino acid modification	1120	3.3e-10
7	GO:CC	GO:0015630	microtubule cytoskeleton	1271	1.7e-08
8	GO:CC	GO:0005815	microtubule organizing center	787	7.1e-08
9	REAC	REAC:R-HSA-1430728	Metabolism	1832	2.5e-07
10	GO:CC	GO:0005794	Golgi apparatus	1475	6.6e-07
11	KEGG	KEGG:01100	Metabolic pathways	1334	8.6e-07
12	KEGG	KEGG:01522	Endocrine resistance	92	9.1e-07
13	KEGG	KEGG:04072	Phospholipase D signaling pathway	138	5.8e-06
14	REAC	REAC:R-HSA-556833	Metabolism of lipids	648	9.5e-06
15	REAC	REAC:R-HSA-199991	Membrane Trafficking	578	1.8e-04

g:Profiler (biit.cs.ut.ee/gprofiler)