



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0000166	nucleotide binding	1973	4.2e-13
2	GO:MF	GO:1901265	nucleoside phosphate binding	1974	4.3e-13
3	GO:MF	GO:0005524	ATP binding	1369	1.1e-11
4	GO:BP	GO:0006259	DNA metabolic process	973	4.6e-09
5	GO:BP	GO:0006281	DNA repair	559	5.2e-08
6	GO:CC	GO:0005794	Golgi apparatus	1475	4.7e-07
7	GO:CC	GO:0005815	microtubule organizing center	787	5.0e-07
8	GO:BP	GO:0033043	regulation of organelle organization	1106	5.2e-07
9	GO:CC	GO:0005813	centrosome	596	1.9e-06
10	REAC	REAC:R-HSA-1483257	Phospholipid metabolism	191	2.5e-05
11	REAC	REAC:R-HSA-556833	Metabolism of lipids	648	2.6e-05
12	REAC	REAC:R-HSA-199991	Membrane Trafficking	578	7.5e-05
13	KEGG	KEGG:01100	Metabolic pathways	1334	7.6e-05
14	KEGG	KEGG:00630	Glyoxylate and dicarboxylate metabolism	28	1.9e-02
15	KEGG	KEGG:04380	Osteoclast differentiation	111	2.2e-02

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