



id	source	term_id	term_name	term_size	p_valu
1	REAC	REAC:R-HSA-1640170	Cell Cycle	640	1.9e-16
2	REAC	REAC:R-HSA-69278	Cell Cycle, Mitotic	517	5.5e-14
3	GO:BP	GO:0022402	cell cycle process	1173	2.3e-13
4	GO:BP	GO:0007049	cell cycle	1655	5.1e–13
5	GO:CC	GO:0005694	chromosome	1638	3.3e-13
6	GO:BP	GO:0000278	mitotic cell cycle	878	3.7e-12
7	REAC	REAC:R-HSA-69620	Cell Cycle Checkpoints	275	1.5e-9
8	GO:CC	GO:0098687	chromosomal region	366	2.5e-7
9	GO:CC	GO:0000785	chromatin	1112	3.4e-6
10	GO:MF	GO:0030527	structural constituent of chromatin	78	3.9e-5
11	KEGG	KEGG:04110	Cell cycle	123	2.1e-4
12	KEGG	KEGG:05322	Systemic lupus erythematosus	101	3.2e-4
13	KEGG	KEGG:04613	Neutrophil extracellular trap formation	149	1.4e-3
14	GO:MF	GO:0140097	catalytic activity, acting on DNA	223	1.9e-2
15	GO:MF	GO:0046982	protein heterodimerization activity	293	5.9e-2

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