



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-1640170	Cell Cycle	640	7.0e-157
2	REAC	REAC:R-HSA-69278	Cell Cycle, Mitotic	517	6.3e-146
3	GO:BP	GO:0007049	cell cycle	1655	6.8e-141
4	GO:BP	GO:0022402	cell cycle process	1173	2.4e-138
5	GO:CC	GO:0005694	chromosome	1638	1.1e-129
6	GO:BP	GO:0000278	mitotic cell cycle	878	1.5e-121
7	REAC	REAC:R-HSA-69620	Cell Cycle Checkpoints	275	1.0e-91
8	GO:CC	GO:0098687	chromosomal region	366	1.0e-77
9	GO:CC	GO:0000775	chromosome, centromeric region	235	7.0e-62
10	GO:MF	GO:0030527	structural constituent of chromatin	78	1.7e-55
11	KEGG	KEGG:04110	Cell cycle	123	3.2e-40
12	KEGG	KEGG:05322	Systemic lupus erythematosus	101	1.5e-39
13	KEGG	KEGG:05034	Alcoholism	154	5.5e-33
14	GO:MF	GO:0140097	catalytic activity, acting on DNA	223	3.5e-27
15	GO:MF	GO:0003682	chromatin binding	518	1.8e-25

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