



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-1640170	Cell Cycle	640	1.8e-166
2	REAC	REAC:R-HSA-69278	Cell Cycle, Mitotic	517	1.5e-154
3	GO:CC	GO:0005694	chromosome	1638	2.8e-145
4	GO:BP	GO:0007049	cell cycle	1655	9.4e-142
5	GO:BP	GO:0022402	cell cycle process	1173	9.0e-139
6	GO:BP	GO:0000278	mitotic cell cycle	878	1.9e-123
7	REAC	REAC:R-HSA-69620	Cell Cycle Checkpoints	275	6.1e-97
8	GO:CC	GO:0098687	chromosomal region	366	4.5e-84
9	GO:CC	GO:0000775	chromosome, centromeric region	235	1.4e-70
10	GO:MF	GO:0030527	structural constituent of chromatin	78	4.3e-57
11	KEGG	KEGG:04110	Cell cycle	123	5.2e-43
12	KEGG	KEGG:05322	Systemic lupus erythematosus	101	9.2e-41
13	KEGG	KEGG:05034	Alcoholism	154	8.0e-35
14	GO:MF	GO:0140097	catalytic activity, acting on DNA	223	1.5e-33
15	GO:MF	GO:0003682	chromatin binding	518	2.2e-31

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