



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0045202	synapse	1219	5.7e-14
2	GO:CC	GO:0030054	cell junction	1916	7.2e-11
3	GO:CC	GO:0043005	neuron projection	1254	2.2e-09
4	GO:BP	GO:0051049	regulation of transport	1545	4.9e-07
5	REAC	REAC:R-HSA-112316	Neuronal System	349	7.9e-06
6	GO:BP	GO:0032879	regulation of localization	1863	9.5e-06
7	GO:BP	GO:0048699	generation of neurons	1343	4.6e-05
8	KEGG	KEGG:04015	Rap1 signaling pathway	199	4.8e-04
9	REAC	REAC:R-HSA-392517	Rap1 signalling	14	9.4e-04
10	GO:MF	GO:0030695	GTPase regulator activity	461	3.8e-03
11	GO:MF	GO:0060589	nucleoside-triphosphatase regulator activity	461	3.8e-03
12	REAC	REAC:R-HSA-6794362	Protein-protein interactions at synapses	76	5.6e-03
13	GO:MF	GO:0098772	molecular function regulator activity	1651	8.2e-03
14	KEGG	KEGG:04911	Insulin secretion	75	1.2e-02
15	KEGG	KEGG:04261	Adrenergic signaling in cardiomyocytes	141	1.6e-02

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