



id	source	term_id	torm name	term_size		p_value
Iu	+	terni_id	term_name	term_size		p_value
1	GO:CC	GO:0043005	neuron projection	1254	4	4.7e–33
2	GO:CC	GO:0045202	synapse	1219	2	2.4e-32
3	GO:CC	GO:0030054	cell junction	1916	6	6.0e-28
4	GO:BP	GO:0120036	plasma membrane bounded cell projection organization	1399	4	4.8e–22
5	GO:BP	GO:0030030	cell projection organization	1436	7	7.0e–22
6	GO:BP	GO:0048666	neuron development	1031	4	4.6e–18
7	GO:MF	GO:0008092	cytoskeletal protein binding	909	4	4.3e-13
8	GO:MF	GO:0019899	enzyme binding	1888	3	3.1e-11
9	GO:MF	GO:0000166	nucleotide binding	1973	3	3.9e-11
10	REAC	REAC:R-HSA-199991	Membrane Trafficking	578	1	1.5e-10
11	REAC	REAC:R-HSA-983168	Antigen processing: Ubiquitination & Proteasome degradation	285	3	3.8e-09
12	REAC	REAC:R-HSA-112316	Neuronal System	349	7	7.1e-09
13	KEGG	KEGG:04727	GABAergic synapse	78	-	1.2e-08
14	KEGG	KEGG:01100	Metabolic pathways	1334	-	1.8e-07
15	KEGG	KEGG:04150	mTOR signaling pathway	144	3	3.5e-07

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