



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0030425	dendrite	587	8.3e-18
2	GO:CC	GO:0097447	dendritic tree	589	9.4e-18
3	GO:CC	GO:0045202	synapse	1219	3.6e-16
4	REAC	REAC:R-HSA-112316	Neuronal System	349	1.2e-09
5	GO:BP	GO:0050804	modulation of chemical synaptic transmission	410	6.4e-09
6	GO:BP	GO:0099177	regulation of trans-synaptic signaling	411	6.8e-09
7	GO:BP	GO:0051049	regulation of transport	1545	1.3e-08
8	REAC	REAC:R-HSA-6794362	Protein-protein interactions at synapses	76	1.8e-07
9	REAC	REAC:R-HSA-6794361	Neurexins and neuroligins	51	7.8e-05
10	KEGG	KEGG:04724	Glutamatergic synapse	107	9.6e-05
11	GO:MF	GO:0019899	enzyme binding	1888	1.0e-04
12	KEGG	KEGG:04921	Oxytocin signaling pathway	145	3.7e-04
13	GO:MF	GO:0140110	transcription regulator activity	1632	9.4e-04
14	GO:MF	GO:0050321	tau-protein kinase activity	22	1.2e-03
15	KEGG	KEGG:04911	Insulin secretion	75	3.9e-03

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