



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0098588	bounding membrane of organelle	1933	1.1e-08
2	GO:CC	GO:0030054	cell junction	1916	4.1e-08
3	GO:CC	GO:0005794	Golgi apparatus	1475	5.5e-08
4	GO:MF	GO:0019899	enzyme binding	1888	3.7e-07
5	GO:BP	GO:0016192	vesicle-mediated transport	1422	4.0e-06
6	GO:BP	GO:0018107	peptidyl-threonine phosphorylation	106	1.3e-05
7	GO:BP	GO:0050804	modulation of chemical synaptic transmission	410	1.7e-05
3	GO:MF	GO:0032555	purine ribonucleotide binding	1739	2.1e-05
9	GO:MF	GO:0032553	ribonucleotide binding	1754	2.5e-05
0	REAC	REAC:R-HSA-112316	Neuronal System	349	3.8e-04
1	KEGG	KEGG:04911	Insulin secretion	75	5.9e-04
2	KEGG	KEGG:04971	Gastric acid secretion	68	1.6e-03
3	KEGG	KEGG:04976	Bile secretion	59	3.5e-03
4	REAC	REAC:R-HSA-5653656	Vesicle-mediated transport	614	7.3e-03
5	REAC	REAC:R-HSA-163685	Integration of energy metabolism	100	1.1e-02
g:Profiler (bijt cs.ut ee/aprofi					orofiler)

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