



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0140098	catalytic activity, acting on RNA	375	2.7e-08
2	GO:MF	GO:0035639	purine ribonucleoside triphosphate binding	1676	1.1e-07
3	GO:MF	GO:0032555	purine ribonucleotide binding	1739	2.9e-07
4	GO:BP	GO:0016570	histone modification	434	1.0e-06
5	GO:CC	GO:0005694	chromosome	1638	2.3e-06
6	GO:CC	GO:0016604	nuclear body	796	5.0e-06
7	GO:BP	GO:0034660	ncRNA metabolic process	586	1.1e-05
8	GO:BP	GO:0070647	protein modification by small protein conjugation or removal	919	1.1e-05
9	GO:CC	GO:0016607	nuclear speck	394	5.3e-05
0	REAC	REAC:R-HSA-1989781	PPARA activates gene expression	105	1.3e-03
1	REAC	REAC:R-HSA-400206	Regulation of lipid metabolism by PPARalpha	106	1.4e-03
12	REAC	REAC:R-HSA-381340	Transcriptional regulation of white adipocyte differentiation	75	1.6e-03
13	KEGG	KEGG:04152	AMPK signaling pathway	109	3.6e-02
14	KEGG	KEGG:04211	Longevity regulating pathway	83	4.6e-02
15	KEGG	KEGG:04910	Insulin signaling pathway	120	4.9e-02

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