IG	Source	term_id	term_name	term_size	intersection_size	p_value
1	GO:BP	GO:0017148	negative regulation of translation	86	7	1.2e-03
2	GO:BP	GO:0006518	peptide metabolic process	575	15	2.2e-03
3	GO:BP	GO:0034249	negative regulation of amide metabolic process	95	7	2.4e-03
4	GO:BP	GO:0043603	amide metabolic process	687	16	4.2e-03
5	GO:BP	GO:0006412	translation	469	12	3.4e-02
6	GO:CC	GO:0005829	cytosol	1105	18	2.1e-02
7	GO:MF	GO:0005488	binding	6905	64	1.4e-05
8	GO:MF	GO:0003727	single-stranded RNA binding	42	4	4.8e-02

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

torm size intersection size