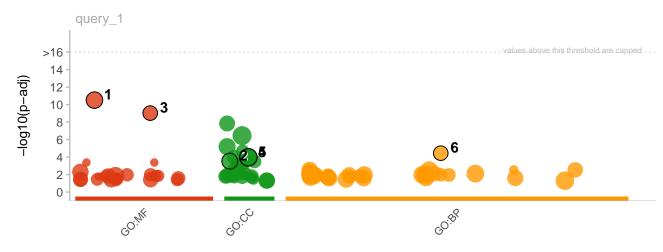
Sr Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	61	3.6e-05
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	502	1.1e-04
4	GO:CC	GO:0043228	non-membrane-bounded organelle	502	1.1e-04
2	GO:CC	GO:0005840	ribosome	155	2.9e-04
1	GO:MF	GO:0005198	structural molecule activity	236	3.1e-11
3	GO:MF	GO:0042302	structural constituent of cuticle	64	9.4e-10

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