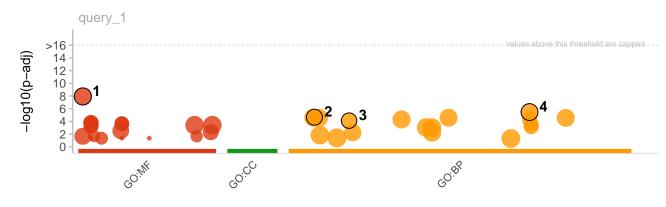
Sp Bottom 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
4	GO:BP	GO:0090304	nucleic acid metabolic process	1218	3.4e-06
2	GO:BP	GO:0006259	DNA metabolic process	469	2.2e-05
3	GO:BP	GO:0015074	DNA integration	297	7.6e-05
1	GO:MF	GO:0003676	nucleic acid binding	1616	1.2e-08

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