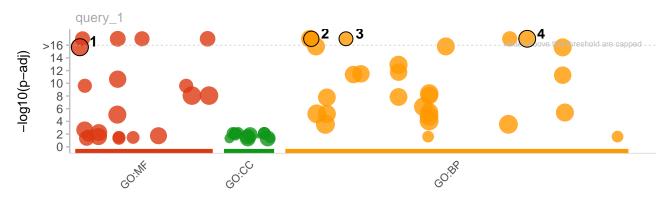
## Sr Bottom 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
2	GO:BP	GO:0006259	DNA metabolic process	204	3.9e-39
3	GO:BP	GO:0015074	DNA integration	60	8.7e-30
4	GO:BP	GO:0090304	nucleic acid metabolic process	909	8.7e-20
1	GO:MF	GO:0003676	nucleic acid binding	1149	2.1e-16

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