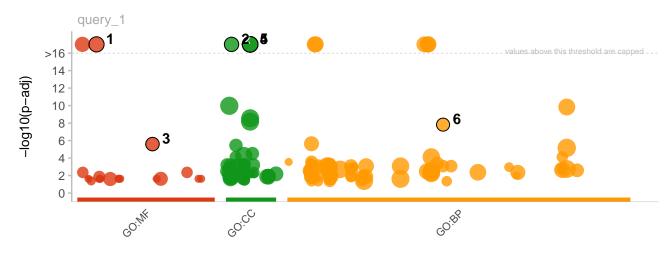
## Sv Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	47	1.5e-08
2	GO:CC	GO:0005840	ribosome	121	1.5e-27
4	GO:CC	GO:0043228	non-membrane-bounded organelle	381	1.5e-24
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	381	1.5e-24
1	GO:MF	GO:0005198	structural molecule activity	236	5.4e-29
3	GO:MF	GO:0042302	structural constituent of cuticle	65	2.5e-06

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