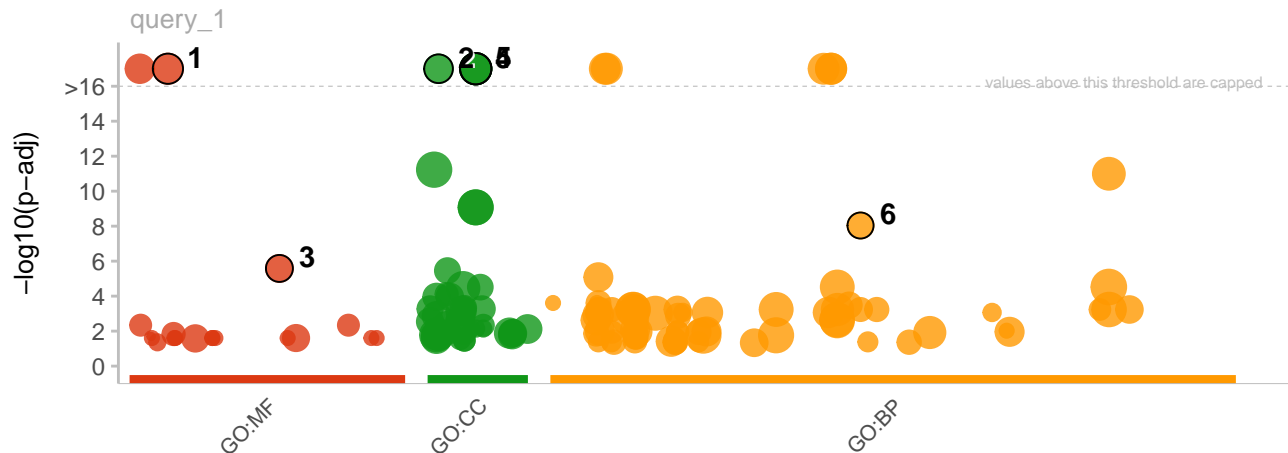


# Sv Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	47	9.2e-09
2	GO:CC	GO:0005840	ribosome	122	5.3e-29
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	383	9.4e-26
4	GO:CC	GO:0043228	non-membrane-bounded organelle	383	9.4e-26
1	GO:MF	GO:0005198	structural molecule activity	230	1.4e-29
3	GO:MF	GO:0042302	structural constituent of cuticle	65	2.6e-06