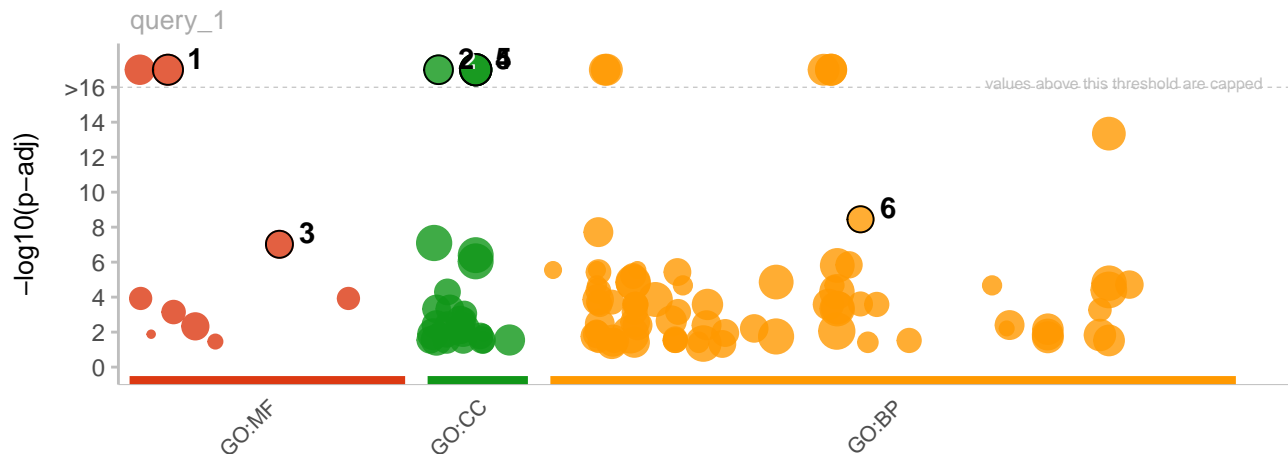


# Sp Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	51	3.6e-09
2	GO:CC	GO:0005840	ribosome	134	2.6e-27
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	417	6.8e-20
4	GO:CC	GO:0043228	non-membrane-bounded organelle	417	6.8e-20
1	GO:MF	GO:0005198	structural molecule activity	243	9.4e-33
3	GO:MF	GO:0042302	structural constituent of cuticle	70	9.5e-08