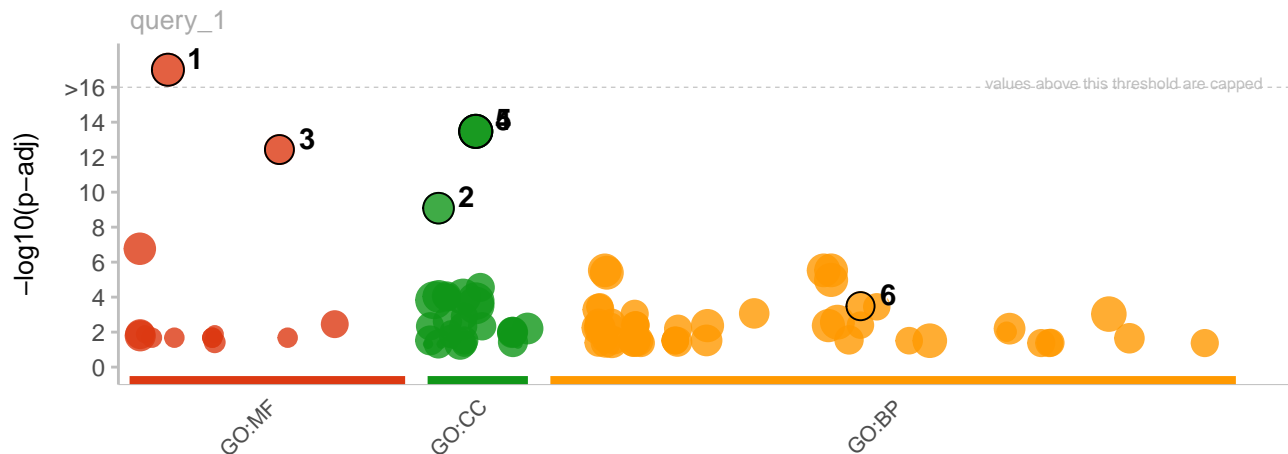


# Ss Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	46	3.3e-04
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	374	3.3e-14
4	GO:CC	GO:0043228	non-membrane-bounded organelle	374	3.3e-14
2	GO:CC	GO:0005840	ribosome	125	8.2e-10
1	GO:MF	GO:0005198	structural molecule activity	231	1.1e-20
3	GO:MF	GO:0042302	structural constituent of cuticle	64	3.7e-13