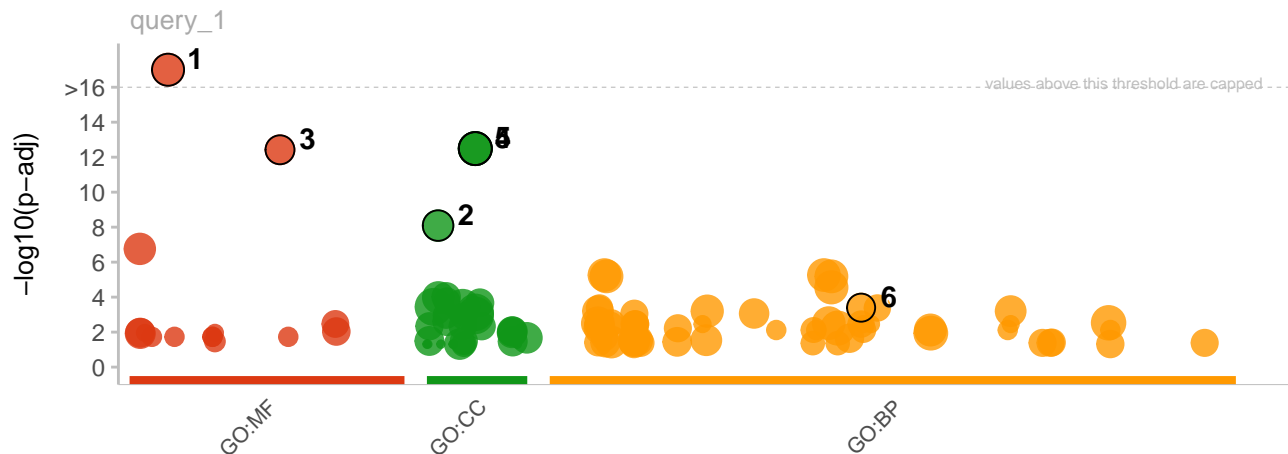


Ss Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	46	4.0e-04
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	370	3.2e-13
4	GO:CC	GO:0043228	non-membrane-bounded organelle	370	3.2e-13
2	GO:CC	GO:0005840	ribosome	123	8.1e-09
1	GO:MF	GO:0005198	structural molecule activity	237	2.5e-21
3	GO:MF	GO:0042302	structural constituent of cuticle	64	3.8e-13