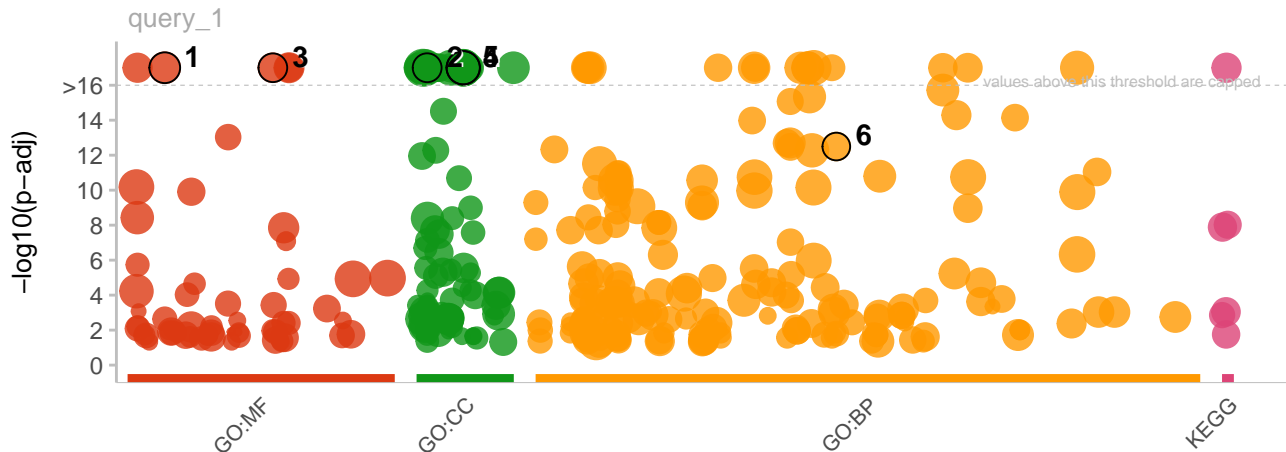


# Ce Top 2 % Codon Adapted Genes



id	source	term_id	term_name	term_size	p_value
6	GO:BP	GO:0046034	ATP metabolic process	109	3.2e-13
4	GO:CC	GO:0043228	non-membrane-bounded organelle	1438	3.3e-82
5	GO:CC	GO:0043232	intracellular non-membrane-bounded organelle	1438	3.3e-82
2	GO:CC	GO:0005840	ribosome	178	1.1e-74
1	GO:MF	GO:0005198	structural molecule activity	407	1.5e-128
3	GO:MF	GO:0042302	structural constituent of cuticle	167	3.3e-56