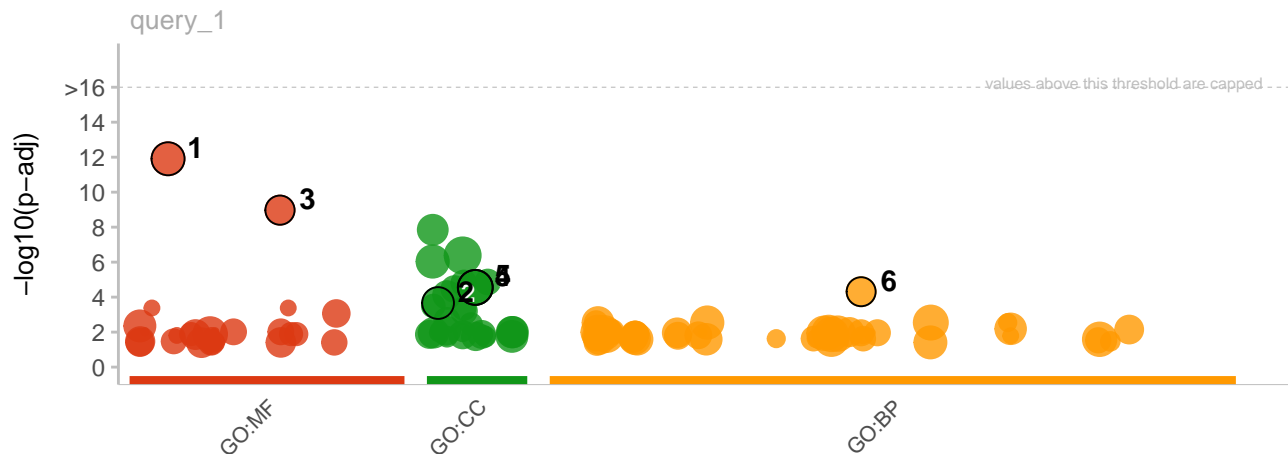


Sr Top 2 % Codon Adapted Genes



| id | source | term_id | term_name | term_size | p_value |
|----|--------|------------|--|-----------|---------|
| 6 | GO:BP | GO:0046034 | ATP metabolic process | 61 | 4.9e-05 |
| 5 | GO:CC | GO:0043232 | intracellular non-membrane-bounded organelle | 502 | 2.8e-05 |
| 4 | GO:CC | GO:0043228 | non-membrane-bounded organelle | 502 | 2.8e-05 |
| 2 | GO:CC | GO:0005840 | ribosome | 155 | 2.2e-04 |
| 1 | GO:MF | GO:0005198 | structural molecule activity | 243 | 1.2e-12 |
| 3 | GO:MF | GO:0042302 | structural constituent of cuticle | 64 | 1.1e-09 |