| id | source | term_id | term_name | term_size | intersection_size | | p_value |
|---|---|---|---|---|---|---|---|
| 1 | GO:BP | GO:0005975 | carbohydrate metabolic process | 963 | 16 | | 4.9e−04 |
| 2 | GO:CC | GO:0048046 | apoplast | 146 | 5 | | 6.8e−03 |
| 3 | GO:MF | GO:0030145 | manganese ion binding | 54 | 4 | | 1.4e−02 |
| 4 | GO:MF | GO:0003824 | catalytic activity | 8918 | 52 | | 2.0e−02 |
| 5 | GO:MF | GO:0045735 | nutrient reservoir activity | 75 | 4 | | 2.0e−02 |
| 6 | GO:MF | GO:0016835 | carbon−oxygen lyase activity | 152 | 5 | | 2.0e−02 |
| 7 | GO:MF | GO:0004096 | catalase activity | 9 | 2 | | 2.1e−02 |
| 8 | GO:CC | GO:0005576 | extracellular region | 487 | 7 | | 2.9e−02 |
| 9 | GO:MF | GO:0016491 | oxidoreductase activity | 1821 | 17 | | 3.1e−02 |
| 10 | GO:MF | GO:0004869 | cysteine−type endopeptidase inhibitor activity | 12 | 2 | | 3.1e−02 |
| 11 | GO:MF | GO:0010333 | terpene synthase activity | 54 | 3 | | 3.3e−02 |
| 12 | GO:MF | GO:0016838 | carbon−oxygen lyase activity, acting on phosphates | 61 | 3 | | 4.2e−02 |
| 13 | GO:BP | GO:0016102 | diterpenoid biosynthetic process | 33 | 3 | | 4.3e−02 |
| 14 | GO:BP | GO:0016101 | diterpenoid metabolic process | 34 | 3 | | 4.3e−02 |
| 15 | GO:MF | GO:0016829 | lyase activity | 336 | 6 | | 4.7e−02 |
| 16 | GO:MF | GO:0008453 | alanine−glyoxylate transaminase activity | 1 | 1 | | 4.7e−02 |
| 17 | GO:MF | GO:0043169 | cation binding | 2798 | 21 | | 4.7e−02 |
| 18 | GO:MF | GO:0004857 | enzyme inhibitor activity | 151 | 4 | | 4.7e−02 |
| 19 | GO:MF | GO:0046872 | metal ion binding | 2785 | 21 | | 4.7e−02 |
| 20 | GO:MF | GO:0016798 | hydrolase activity, acting on glycosyl bonds | 496 | 7 | | 4.7e−02 |
| 21 | GO:MF | GO:0004553 | hydrolase activity, hydrolyzing O−glycosyl compounds | 457 | 7 | | 4.7e−02 |
| 22 | GO:MF | GO:0016174 | NAD(P)H oxidase H2O2−forming activity | 1 | 1 | | 4.7e−02 |
| 23 | GO:MF | GO:0033946 | xyloglucan−specific endo−beta−1,4−glucanase activity | 1 | 1 | | 4.7e−02 |