

| id | source | term_id    | term_name                                    | term_size | intersection_size | p_value |
|----|--------|------------|--|-----------|-------------------|---------|
| 1  | GO:BP  | GO:0001666 | response to hypoxia                          | 266       | 12                | 2.4e-02 |
| 2  | GO:BP  | GO:0036293 | response to decreased oxygen levels          | 270       | 12                | 2.8e-02 |
| 3  | GO:BP  | GO:0070482 | response to oxygen levels                    | 271       | 12                | 2.9e-02 |
| 4  | GO:BP  | GO:0008033 | tRNA processing                              | 115       | 5                 | 3.0e-02 |
| 5  | GO:BP  | GO:0071456 | cellular response to hypoxia                 | 239       | 11                | 4.3e-02 |
| 6  | GO:BP  | GO:0036294 | cellular response to decreased oxygen levels | 241       | 11                | 4.6e-02 |
| 7  | GO:BP  | GO:0071453 | cellular response to oxygen levels           | 241       | 11                | 4.6e-02 |
| 8  | GO:CC  | GO:0071944 | cell periphery                               | 2902      | 59                | 2.9e-04 |
| 9  | GO:CC  | GO:0005886 | plasma membrane                              | 2506      | 50                | 3.8e-03 |
| 10 | GO:CC  | GO:0005911 | cell-cell junction                           | 900       | 18                | 1.3e-02 |
| 11 | GO:CC  | GO:0009506 | plasmodesma                                  | 900       | 18                | 1.3e-02 |
| 12 | GO:CC  | GO:0070161 | anchoring junction                           | 900       | 18                | 1.3e-02 |
| 13 | GO:CC  | GO:0055044 | symplast                                     | 901       | 18                | 1.3e-02 |
| 14 | GO:CC  | GO:0030054 | cell junction                                | 906       | 18                | 1.4e-02 |
| 15 | GO:MF  | GO:0036094 | small molecule binding                       | 6709      | 55                | 3.6e-03 |
| 16 | GO:MF  | GO:0043167 | ion binding                                  | 6591      | 54                | 4.7e-03 |
| 17 | GO:MF  | GO:0003972 | RNA ligase (ATP) activity                    | 1         | 1                 | 2.5e-02 |
| 18 | GO:MF  | GO:0008452 | RNA ligase activity                          | 1         | 1                 | 2.5e-02 |
| 19 | GO:MF  | GO:0033612 | receptor serine/threonine kinase binding     | 75        | 2                 | 3.0e-02 |