



| id | source | term_id            | term_name                                       | term_size | p_value |
|----|--------|--------------------|---|-----------|---------|
| 1  | REAC   | REAC:R-HSA-8953854 | Metabolism of RNA                               | 626       | 1.9e-96 |
| 2  | GO:MF  | GO:0003723         | RNA binding                                     | 1885      | 1.6e-71 |
| 3  | REAC   | REAC:R-HSA-74160   | Gene expression (Transcription)                 | 1380      | 1.4e-63 |
| 4  | REAC   | REAC:R-HSA-72203   | Processing of Capped Intron–Containing Pre–mRNA | 226       | 2.8e-60 |
| 5  | GO:CC  | GO:1902494         | catalytic complex                               | 1628      | 8.1e-58 |
| 6  | GO:CC  | GO:0005681         | spliceosomal complex                            | 184       | 4.3e-45 |
| 7  | GO:BP  | GO:0034660         | ncRNA metabolic process                         | 586       | 2.3e-43 |
| 8  | GO:CC  | GO:0016604         | nuclear body                                    | 796       | 2.8e-42 |
| 9  | GO:BP  | GO:0006259         | DNA metabolic process                           | 973       | 1.8e-39 |
| 10 | GO:BP  | GO:0042254         | ribosome biogenesis                             | 293       | 4.9e-38 |
| 11 | KEGG   | KEGG:03040         | Spliceosome                                     | 133       | 8.2e-32 |
| 12 | GO:MF  | GO:0140640         | catalytic activity, acting on a nucleic acid    | 574       | 2.0e-30 |
| 13 | GO:MF  | GO:0140098         | catalytic activity, acting on RNA               | 375       | 8.6e-26 |
| 14 | KEGG   | KEGG:03013         | Nucleocytoplasmic transport                     | 98        | 8.1e-16 |
| 15 | KEGG   | KEGG:03008         | Ribosome biogenesis in eukaryotes               | 72        | 4.4e-13 |

g:Profiler (biit.cs.ut.ee/gprofiler)