```
116/432 cellular component assembly
10/12 ribosomal large subunit assembly
86/198 cellular macromolecular complex assembly
4/67 ribosomal small subunit assembly
105/355 macromolecular complex subunit organization
8/9 ribosomal small subunit assembly
105/355 macromolecular complex subunit organization
29/67 protein—DNA complex subunit organization
92/269 macromolecular complex assembly
49/191 protein complex assembly
                                                                                                                                                                                                                             p < 0.01 p < 0.05
 49/191 protein complex assembly
49/191 protein complex assembly
127/575 organelle organization
26/56 nucleosome organization
67/207 chromosome organization
114/193 translation
67/ formation of translation preinitiation complex
16/31 translational initiation
22/716 macromolecule biosynthetic process
7/9 cytoplasmic translation on translation 10/15 regulation of translational initiation
25/54 regulation of translation
25/54 regulation of gene expression
32/62 embryo development
13/33 gene silencing
11/19 chromatin silencing
        34 regulation of gene expression, epigenetic
103/440 response to stress
50/186 cellular response to DNA damage stimulus
 58/220 DNA metabolic process
31/86 protein folding
48/176 regulation of cell cycle
 23/68 regulation of cell cycle process 36/104 cell cycle
6/7 rRNA 3'-end processing
10/21 RNA 3'-end processing
8/10 ncRNA 3'-end processing
41/76 rRNA metabolic process
57 maturation of LSU-rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, LSU 67/162 ncRNA metabolic process 8/11 RNA phosphodiester bond hydrolysis, endonucleolytic 14/35 nucleic acid phosphodiester bond hydrolysis 5/5 endonucleolytic cleavage of tricistronic rRNA transcript (SSU-rR 9/10 cleavage involved in rRNA processing 226/767 RNA metabolic process 13/18 RNA phosphodiester bond hydrolysis 123/280 RNA processing 34/54 RNA splicing, via transesterification reactions 60/141 mRNA metabolic process 21/58 RNA splicing 1/58 RNA modification
                                                                                            tronic rRNA transcript (SSU-rRNA, 5.8S rRNA, LSU
21/58 RNA modification 7/10 pseudouridine synthesis
26/81 tRNA metabolic process
19/31 ribosome biogenesis
26/44 cellular component biogenesis
60/196 macromolecule catabolic process
25/73 protein catabolic process
55/157 proteolysis involved in cellular protein catabolic process
10/18 mRNA transport
14/33 nucleobase—containing compound transport 7/10 regulation of endodermal cell fate specification 14/25 electron transport chain
 4/5 mitochondrial electron transport, ubiquinol to cytochrome c
 6/10 aerobic respiration
 6/10 ketone biosynthetic process
 4/5 polyprenol metabolic process
```