# GO term of Cluster2

RNA processing
ncRNA metabolic process
ribonucleoprotein complex biogenesis
mRNA metabolic process
ncRNA processing
ribosome biogenesis
mRNA processing
rRNA metabolic process
rRNA processing
animal organ morphogenesis
G protein−coupled receptor signaling pathway
ribonucleoprotein complex subunit organization
ribonucleoprotein complex assembly
regulation of small GTPase mediated signal transduction
positive regulation of GTPase activity
protein complex oligomerization
embryonic organ development
ribosomal small subunit biogenesis
ribosomal large subunit biogenesis
protein homooligomerization
maturation of SSU−rRNA
sensory organ morphogenesis
maturation of SSU−rRNA from tricistronic rRNA transcript (SSU−rRNA, 5.8S rRNA, LSU−rRNA)

BP

cell projection
plasma membrane bounded cell projection
nucleolus
ribonucleoprotein complex
neuron projection
plasma membrane region
intrinsic component of plasma membrane
somatodendritic compartment
integral component of plasma membrane
dendrite
dendritic tree
cell body
spliceosomal complex
neuronal cell body
preribosome
plasma membrane protein complex
cell projection membrane
U2−type spliceosomal complex
catalytic step 2 spliceosome
neuron spine
dendritic spine
cluster of actin−based cell projections
small−subunit processome
RNA polymerase II transcription regulator complex
90S preribosome
sarcolemma
preribosome, large subunit precursor
SWI/SNF superfamily−type complex
neuromuscular junction
GTPase complex
heterotrimeric G−protein complex
preribosome, small subunit precursor
eukaryotic 43S preinitiation complex
translation preinitiation complex

CC

hydrolase activity, acting on acid anhydrides
hydrolase activity, acting on acid anhydrides, in phosphorus−containing anhydrides
pyrophosphatase activity
ribonucleoside triphosphate phosphatase activity
catalytic activity, acting on RNA
GTPase activator activity
ATP−dependent activity, acting on RNA
RNA helicase activity

MF

0.00
0.05
0.10
0.15
GeneRatio

Count
100
200

p.adjust
0.04
0.03
0.02
0.01