

ribosome biogenesis, GO:0042254

fibrillar center, GO:0001650
ribosome binding, GO:0043022
preribosome, GO:0030687
snoRNA binding, GO:0030515
pseudouridine synthesis, GO:0001522
positive regulation of interferon-beta production, GO:0032728
ribosomal small subunit binding, GO:0043024
positive regulation of telomerase RNA localization to Cajal body, GO:1904874
pre-miRNA processing, GO:0031054
ribosomal large subunit binding, GO:0043023
positive regulation of establishment of protein localization to telomere, GO:1904851
negative regulation of protein complex assembly, GO:0031333
positive regulation of protein sumoylation, GO:0033235
regulation of protein catabolic process, GO:0042176
nuclear export signal receptor activity, GO:0005049
maturation of 5.8S rRNA, GO:0000460
endoribonuclease activity, GO:0004521
preribosome, GO:0030688
negative regulation of protein sumoylation, GO:0033234
negative regulation of protein homooligomerization, GO:0032463
ribosomal large subunit export from nucleus, GO:0000055
negative regulation of phosphatidylinositol 3-kinase signaling, GO:0014067
GTP metabolic process, GO:0046039
rRNA catabolic process, GO:0016075
negative regulation of reactive oxygen species metabolic process, GO:2000378
rRNA transcription, GO:0009303
homeostasis of number of cells, GO:0048872
T cell mediated immunity, GO:0002456
annulate lamellae, GO:0005642
ribosome assembly, GO:0042255
regulation of regulatory T cell differentiation, GO:0045589
blood coagulation, GO:0007597
protein localization to cytoplasmic stress granule, GO:1903608
histone H3-K4 demethylation, GO:0034720
negative regulation of RIG-I signaling pathway, GO:0039536
RNA polymerase I core binding, GO:0001042
regulation of protein export from nucleus, GO:0046825
production of siRNA involved in RNA interference, GO:0030422
cellular response to arsenic-containing substance, GO:0071243
positive regulation of rRNA processing, GO:2000234
histone demethylase activity (H3-K4 specific), GO:0032453
DEAD/H-box RNA helicase binding, GO:0017151