

ribosome biogenesis, GO:0042254

- fibrillar center, GO:0001650
- small-subunit processome, GO:0032040
- snoRNA binding, GO:0030515
- preribosome, GO:0030687
- negative regulation of protein ubiquitination, GO:0031397
- positive regulation of telomerase RNA localization to Cajal body, GO:1904874
- maturation of 5.8S rRNA, GO:0000460
- positive regulation of interferon-beta production, GO:0032728
- pseudouridine synthesis, GO:0001522
- positive regulation of establishment of protein localization to telomere, GO:1904851
- stress granule assembly, GO:0034063
- RNA modification, GO:0009451
- ribosome assembly, GO:0042255
- nuclear export signal receptor activity, GO:0005049
- histone demethylase activity (H3-K36 specific), GO:0051864
- histone H3-K36 demethylation, GO:0070544
- RNA polymerase binding, GO:0070063
- regulation of centrosome duplication, GO:0010824
- negative regulation of protein complex assembly, GO:0031333
- preribosome, GO:0030688
- pseudouridine synthase activity, GO:0009982
- endonucleolytic cleavage in 5'-ETS of tricistronic rRNA transcript (SSU-rRNA, GO:0000480
- RNA polymerase I CORE element sequence-specific DNA binding, GO:0001164
- ribosomal large subunit export from nucleus, GO:0000055
- cellular response to virus, GO:0098586
- protein localization to cytoplasmic stress granule, GO:1903608
- T cell mediated immunity, GO:0002456
- negative regulation of protein sumoylation, GO:0033234
- endoribonuclease activity, GO:0004521
- positive regulation of protein homodimerization activity, GO:0090073
- negative regulation of reactive oxygen species metabolic process, GO:2000378
- negative regulation of protein homooligomerization, GO:0032463
- Pwp2p-containing subcomplex of 90S preribosome, GO:0034388
- negative regulation of phosphatidylinositol 3-kinase signaling, GO:0014067
- positive regulation of neutrophil chemotaxis, GO:0090023
- cellular response to arsenic-containing substance, GO:0071243
- GTP metabolic process, GO:0046039
- histone H3-K4 demethylation, GO:0034720
- rRNA catabolic process, GO:0016075
- pre-snoRNP complex, GO:0070761
- histone demethylase activity (H3-K4 specific), GO:0032453
- blastocyst formation, GO:0001825
- caspase binding, GO:0089720
- positive regulation of pri-miRNA transcription from RNA polymerase II promoter, GO:1902895
- positive regulation of transcription from RNA polymerase I promoter, GO:0045943
- negative regulation of interleukin-12 production, GO:0032695
- positive regulation of rRNA processing, GO:2000234