

ribosome biogenesis, GO:0042254

- fibrillar center, GO:0001650
- small-subunit processome, GO:0032040
- ATPase binding, GO:0051117
- preribosome, GO:0030687
- snoRNA binding, GO:0030515
- positive regulation of interferon-beta production, GO:0032728
- positive regulation of telomerase RNA localization to Cajal body, GO:1904874
- ribonucleoprotein granule, GO:0035770
- negative regulation of protein complex assembly, GO:0031333
- maturation of 5.8S rRNA, GO:0000460
- RNA stem-loop binding, GO:0035613
- pseudouridine synthesis, GO:0001522
- positive regulation of protein sumoylation, GO:0033235
- T cell mediated immunity, GO:0002456
- RNA polymerase binding, GO:0070063
- trophoblast cell differentiation, GO:0001829
- endoribonuclease activity, GO:0004521
- positive regulation of transcription from RNA polymerase I promoter, GO:0045943
- stem cell division, GO:0017145
- preribosome, GO:0030688
- positive regulation of translational initiation, GO:0045948
- nuclear export signal receptor activity, GO:0005049
- protein localization to organelle, GO:0033365
- endonucleolytic cleavage in 5'-ETS of tricistronic rRNA transcript (SSU-rRNA, GO:0000480
- cellular response to zinc ion, GO:0071294
- cellular response to arsenic-containing substance, GO:0071243
- Pwp2p-containing subcomplex of 90S preribosome, GO:0034388
- negative regulation of reactive oxygen species metabolic process, GO:2000378
- nucleoside-triphosphatase activity, GO:0017111
- negative regulation of RIG-I signaling pathway, GO:0039536
- cellular response to osmotic stress, GO:0071470
- protein localization to cytoplasmic stress granule, GO:1903608
- regulation of aerobic respiration, GO:1903715
- regulation of protein export from nucleus, GO:0046825