

ribosome biogenesis, GO:0042254

- fibrillar center, GO:0001650
- small-subunit processome, GO:0032040
- ATPase binding, GO:0051117
- snoRNA binding, GO:0030515
- mitochondrial translation, GO:0032543
- preribosome, GO:0030687
- endoribonuclease activity, GO:0004521
- negative regulation of protein complex assembly, GO:0031333
- maturation of 5.8S rRNA, GO:0000460
- pseudouridine synthesis, GO:0001522
- ribonucleoprotein granule, GO:0035770
- RNA stem-loop binding, GO:0035613
- leukocyte chemotaxis, GO:0030595
- RNA modification, GO:0009451
- negative regulation of protein sumoylation, GO:0033234
- rRNA transcription, GO:0009303
- positive regulation of transcription from RNA polymerase I promoter, GO:0045943
- histone H3-K4 demethylation, GO:0034720
- protein localization to organelle, GO:0033365
- ribosomal large subunit export from nucleus, GO:0000055
- trophectodermal cell differentiation, GO:0001829
- endonucleolytic cleavage in 5'-ETS of tricistronic rRNA transcript (SSU-rRNA, GO:0000480
- pseudouridine synthase activity, GO:0009982
- blastocyst formation, GO:0001825