GO_TRNA_PROCESSING, GO_TRNA_PROCESSING

GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY, GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY

GO_NUCLEASE_ACTIVITY, GO_NUCLEASE_ACTIVITY

GO_RNA_METHYLATION, GO_RNA_METHYLATION

GO_RNA_MODIFICATION, GO_RNA_MODIFICATION

GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS

GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY, GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY

GO_AMINO_ACID_ACTIVATION, GO_AMINO_ACID_ACTIVATION

GO_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA, GO_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA

GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS

GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION

GO_LIGASE_ACTIVITY, GO_LIGASE_ACTIVITY

GO_RNA_METHYLTRANSFERASE_ACTIVITY, GO_RNA_METHYLTRANSFERASE_ACTIVITY

GO_TRNA_MODIFICATION, GO_TRNA_MODIFICATION

GO_MATURATION_OF_5_8S_RRNA, GO_MATURATION_OF_5_8S_RRNA

GO_RIBONUCLEASE_ACTIVITY, GO_RIBONUCLEASE_ACTIVITY

GO_3_5_EXONUCLEASE_ACTIVITY, GO_3_5_EXONUCLEASE_ACTIVITY

GO_MRNA_CLEAVAGE_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_FACTOR_COMPLEX

GO_EXONUCLEASE_ACTIVITY, GO_EXONUCLEASE_ACTIVITY

TABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS — GO_ENDONUCLEASE_ACTIVITY, GO_ENDONUCLEASE_ACTIVITY

GO_REGULATION_OF_TRANSLATIONAL_ELONGATION, GO_REGULATION_OF_TRANSLATIONAL_ELONGATION

GO_TRNA_METHYLTRANSFERASE_ACTIVITY, GO_TRNA_METHYLTRANSFERASE_ACTIVITY

GO_MRNA_METHYLATION, GO_MRNA_METHYLATION

GO_EXORIBONUCLEASE_COMPLEX, GO_EXORIBONUCLEASE_COMPLEX

GO_MRNA_MODIFICATION, GO_MRNA_MODIFICATION

GO_NUCLEAR_EXOSOME_RNASE_COMPLEX, GO_NUCLEAR_EXOSOME_RNASE_COMPLEX

GO_TRNA_METHYLATION, GO_TRNA_METHYLATION

GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC

GO_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS

GO_RNA_5_END_PROCESSING, GO_RNA_5_END_PROCESSING

GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_AND_POLYADENYLATION_SPECIFICITY_FACTOR_COMPLEX

GO_CYTOPLASMIC_TRANSLATIONAL_ELONGATION, GO_CYTOPLASMIC_TRANSLATIONAL_ELONGATION

GO_TRNA_5_END_PROCESSING, GO_TRNA_5_END_PROCESSING

GO_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GO_MATURATION_OF_5_8S_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA

GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING, GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING

GO_AMINOACYL_TRNA_SYNTHETASE_MULTIENZYME_COMPLEX, GO_AMINOACYL_TRNA_SYNTHETASE_MULTIENZYME_COMPLEX

GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS

GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5