

CATABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS

- GO_MITOCHONDRIAL_GENE_EXPRESSION, GO_MITOCHONDRIAL_GENE_EXPRESSION
- GO_MITOCHONDRIAL_TRANSLATION, GO_MITOCHONDRIAL_TRANSLATION
- GO_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA, GO_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA
- GO_TRNA_PROCESSING, GO_TRNA_PROCESSING
- GO_TRNA_MODIFICATION, GO_TRNA_MODIFICATION
- GO_RNA_MODIFICATION, GO_RNA_MODIFICATION
- GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY, GO_S_ADENOSYLMETHIONINE_DEPENDENT_METHYLTRANSFERASE_ACTIVITY
- GO_RNA_METHYLATION, GO_RNA_METHYLATION
- GO_RNA_METHYLTRANSFERASE_ACTIVITY, GO_RNA_METHYLTRANSFERASE_ACTIVITY
- GO_NUCLEASE_ACTIVITY, GO_NUCLEASE_ACTIVITY
- GO_TRNA_METHYLTRANSFERASE_ACTIVITY, GO_TRNA_METHYLTRANSFERASE_ACTIVITY
- GO_TRNA_METHYLATION, GO_TRNA_METHYLATION
- GO_EXONUCLEASE_ACTIVITY, GO_EXONUCLEASE_ACTIVITY
- GO_LIGASE_ACTIVITY, GO_LIGASE_ACTIVITY
- GO_AMINO_ACID_ACTIVATION, GO_AMINO_ACID_ACTIVATION
- GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY, GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY
- GO_MATURATION_OF_5_8S_RRNA, GO_MATURATION_OF_5_8S_RRNA
- GO_METAL_CLUSTER_BINDING, GO_METAL_CLUSTER_BINDING
- GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS
- GO_TRNA_BINDING, GO_TRNA_BINDING
- 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_3_5_EXONUCLEASE_ACTIVITY, GO_3_5_EXONUCLEASE_ACTIVITY
- 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_P
- GO_4_IRON_4_SULFUR_CLUSTER_BINDING, GO_4_IRON_4_SULFUR_CLUSTER_BINDING
- GO_RRNA_MODIFICATION, GO_RRNA_MODIFICATION
- GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC
- GO_NCRNA_CATABOLIC_PROCESS, GO_NCRNA_CATABOLIC_PROCESS
- GO_TRNA_WOBBLE_URIDINE_MODIFICATION, GO_TRNA_WOBBLE_URIDINE_MODIFICATION
- GO_NUCLEOID, GO_NUCLEOID
- GO_RIBONUCLEASE_ACTIVITY, GO_RIBONUCLEASE_ACTIVITY
- GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS
- GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY
- GO_TRNA_WOBBLE_BASE_MODIFICATION, GO_TRNA_WOBBLE_BASE_MODIFICATION
- GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_P
- GO_ENDONUCLEASE_ACTIVITY, GO_ENDONUCLEASE_ACTIVITY
- GO_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION, GO_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION
- GO_EXORIBONUCLEASE_ACTIVITY, GO_EXORIBONUCLEASE_ACTIVITY
- GO_GLYCOSYL_COMPOUND_BIOSYNTHETIC_PROCESS, GO_GLYCOSYL_COMPOUND_BIOSYNTHETIC_PROCESS
- GO_TRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY, GO_TRNA_GUANINE_METHYLTRANSFERASE_ACTIVITY
- GO_CARBON_NITROGEN_LIGASE_ACTIVITY_WITH_GLUTAMINE_AS_AMIDO_N_DONOR, GO_CARBON_NITROGEN_LIGASE_ACTIVITY_WITH_GLUTAMINE_AS_AMIDO_N_DONOR
- GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_ENDONUCLEOLYTIC
- GO_EXORIBONUCLEASE_COMPLEX, GO_EXORIBONUCLEASE_COMPLEX
- GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING, GO_CLEAVAGE_INVOLVED_IN_RRNA_PROCESSING
- GO_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS
- GO_AMINOACYL_TRNA_SYNTHETASE_MULTTIENZYME_COMPLEX, GO_AMINOACYL_TRNA_SYNTHETASE_MULTTIENZYME_COMPLEX
- GO_CYTOPLASMIC_EXOSOME_RNASE_COMPLEX, GO_CYTOPLASMIC_EXOSOME_RNASE_COMPLEX
- GO_RRNA_METHYLATION, GO_RRNA_METHYLATION
- GO_NUCLEAR_EXOSOME_RNASE_COMPLEX, GO_NUCLEAR_EXOSOME_RNASE_COMPLEX
- GO_ENDORIBONUCLEASE_COMPLEX, GO_ENDORIBONUCLEASE_COMPLEX
- GO_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION, GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION
- GO_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS, GO_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS
- GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
- GO_TRNA_AMINOACYLATION_FOR_MITOCHONDRIAL_PROTEIN_TRANSLATION, GO_TRNA_AMINOACYLATION_FOR_MITOCHONDRIAL_PROTEIN_TRANSLATION
- GO_POLYADENYLATION_DEPENDENT_RNA_CATABOLIC_PROCESS, GO_POLYADENYLATION_DEPENDENT_RNA_CATABOLIC_PROCESS
- GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE, GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERASE
- GO_ENDORIBONUCLEASE_ACTIVITY, GO_ENDORIBONUCLEASE_ACTIVITY
- GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS
- GO_SNRNA_METABOLIC_PROCESS, GO_SNRNA_METABOLIC_PROCESS
- GO_SNORNA_3_END_PROCESSING, GO_SNORNA_3_END_PROCESSING
- GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC, GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_EXONUCLEOLYTIC
- GO_MITOCHONDRIAL_RNA_PROCESSING, GO_MITOCHONDRIAL_RNA_PROCESSING
- GO_ENDONUCLEASE_COMPLEX, GO_ENDONUCLEASE_COMPLEX
- GO_MITOCHONDRIAL_RNA_MODIFICATION, GO_MITOCHONDRIAL_RNA_MODIFICATION
- GO_RRNA_CATABOLIC_PROCESS, GO_RRNA_CATABOLIC_PROCESS