

AL_GENE_EXPRESSION, GO_MITOCHONDRIAL_GENE_EXPRESSION

- GO_TRNA_METABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS
- GO_MITOCHONDRIAL_TRANSLATION, GO_MITOCHONDRIAL_TRANSLATION
- GO_RIBOSOMAL_SUBUNIT, GO_RIBOSOMAL_SUBUNIT
- GO_TRNA_PROCESSING, GO_TRNA_PROCESSING
- GO_RNA_MODIFICATION, GO_RNA_MODIFICATION
- GO_STRUCTURAL_CONSTITUENT_OF_RIBOSOME, GO_STRUCTURAL_CONSTITUENT_OF_RIBOSOME
- GO_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA, GO_CATALYTIC_ACTIVITY_ACTING_ON_A_TRNA
- GO_TRANSLATIONAL_ELONGATION, GO_TRANSLATIONAL_ELONGATION
- GO_TRNA_MODIFICATION, GO_TRNA_MODIFICATION
- GO_TRANSLATIONAL_TERMINATION, GO_TRANSLATIONAL_TERMINATION
- GO_LARGE_RIBOSOMAL_SUBUNIT, GO_LARGE_RIBOSOMAL_SUBUNIT
- GO_RIBONUCLEOPROTEIN_COMPLEX_BINDING, GO_RIBONUCLEOPROTEIN_COMPLEX_BINDING
- GO_ORGANELLAR_RIBOSOME, GO_ORGANELLAR_RIBOSOME
- GO_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION, GO_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION
- GO_RNA_METHYLATION, GO_RNA_METHYLATION
- GO_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT, GO_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT
- GO_NUCLEOID, GO_NUCLEOID
- GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY, GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY
- GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS
- GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS
- GO_AMINO_ACID_ACTIVATION, GO_AMINO_ACID_ACTIVATION
- GO_SMALL_RIBOSOMAL_SUBUNIT, GO_SMALL_RIBOSOMAL_SUBUNIT
- GO_RNA_METHYLTRANSFERASE_ACTIVITY, GO_RNA_METHYLTRANSFERASE_ACTIVITY
- GO_RIBOSOME_BINDING, GO_RIBOSOME_BINDING
- GO_TRANSLATION_REGULATOR_ACTIVITY, GO_TRANSLATION_REGULATOR_ACTIVITY
- GO_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING, GO_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING
- GO_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS, GO_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS
- GO_LIGASE_ACTIVITY, GO_LIGASE_ACTIVITY
- GO_RRNA_MODIFICATION, GO_RRNA_MODIFICATION
- GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION, GO_POSITIVE_REGULATION_OF_MITOCHONDRIAL_TRANSLATION
- GO_TRNA_WOBBLE_BASE_MODIFICATION, GO_TRNA_WOBBLE_BASE_MODIFICATION
- GO_TRNA_BINDING, GO_TRNA_BINDING
- GO_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION, GO_REGULATION_OF_MITOCHONDRIAL_GENE_EXPRESSION
- GO_PSEUDOURIDINE_SYNTHESIS, GO_PSEUDOURIDINE_SYNTHESIS
- GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY, GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY
- GO_TRNA_METHYLATION, GO_TRNA_METHYLATION
- GO_PSEUDOURIDINE_SYNTHASE_ACTIVITY, GO_PSEUDOURIDINE_SYNTHASE_ACTIVITY
- GO_MITOCHONDRIAL_RNA_PROCESSING, GO_MITOCHONDRIAL_RNA_PROCESSING
- GO_POSITIVE_REGULATION_OF_TRANSLATION, GO_POSITIVE_REGULATION_OF_TRANSLATION
- GO_CYTOPLASMIC_EXOSOME_RNASE_COMPLEX, GO_CYTOPLASMIC_EXOSOME_RNASE_COMPLEX
- GO_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY, GO_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY
- GO_DNA_DAMAGE_RESPONSE_DETECTION_OF_DNA_DAMAGE, GO_DNA_DAMAGE_RESPONSE_DETECTION_OF_DNA_DAMAGE
- GO_TRNA_METHYLTRANSFERASE_ACTIVITY, GO_TRNA_METHYLTRANSFERASE_ACTIVITY
- GO_CYTOCHROME_COMPLEX_ASSEMBLY, GO_CYTOCHROME_COMPLEX_ASSEMBLY
- GO_NCRNA_3_END_PROCESSING, GO_NCRNA_3_END_PROCESSING
- GO_TRANSLATION_REGULATOR_ACTIVITY_NUCLEIC_ACID_BINDING, GO_TRANSLATION_REGULATOR_ACTIVITY_NUCLEIC_ACID_BINDING
- GO_TRNA_3_END_PROCESSING, GO_TRNA_3_END_PROCESSING
- GO_RIBOSOMAL_LARGE_SUBUNIT_BINDING, GO_RIBOSOMAL_LARGE_SUBUNIT_BINDING
- GO_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY, GO_RIBOSOMAL_LARGE_SUBUNIT_ASSEMBLY
- GO_RIBOSOME_ASSEMBLY, GO_RIBOSOME_ASSEMBLY
- GO_RESPIRATORY_CHAIN_COMPLEX_IV_ASSEMBLY, GO_RESPIRATORY_CHAIN_COMPLEX_IV_ASSEMBLY
- GO_ORGANELLE_ENVELOPE_LUMEN, GO_ORGANELLE_ENVELOPE_LUMEN
- GO_RRNA_BINDING, GO_RRNA_BINDING
- GO_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT, GO_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT
- GO_MITOCHONDRIAL_TRANSCRIPTION, GO_MITOCHONDRIAL_TRANSCRIPTION
- GO_MITOCHONDRIAL_TRNA_PROCESSING, GO_MITOCHONDRIAL_TRNA_PROCESSING
- GO_RRNA_METHYLATION, GO_RRNA_METHYLATION
- GO_AMINOACYL_TRNA_EDITING_ACTIVITY, GO_AMINOACYL_TRNA_EDITING_ACTIVITY
- GO_AMINOACYL_TRNA_METABOLISM_INVOLVED_IN_TRANSLATIONAL_FIDELITY, GO_AMINOACYL_TRNA_METABOLISM_INVOLVED_IN_TRANSLATIONAL_FIDELITY
- GO_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION, GO_REGULATION_OF_OXIDATIVE_PHOSPHORYLATION
- GO_ENDONUCLEASE_COMPLEX, GO_ENDONUCLEASE_COMPLEX