AL_GENE_EXPRESSION, GO_MITOCHONDRIAL_GENE_EXPRESSION

- GO_TRANSLATIONAL_ELONGATION, GO_TRANSLATIONAL_ELONGATION
- GO_TRANSLATIONAL_TERMINATION, GO_TRANSLATIONAL_TERMINATION
- GO_MITOCHONDRIAL_TRANSLATION, GO_MITOCHONDRIAL_TRANSLATION
- GO_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION, GO_MITOCHONDRIAL_TRANSLATIONAL_TERMINATION
- GO_ORGANELLAR_RIBOSOME, GO_ORGANELLAR_RIBOSOME
- GO_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT, GO_MITOCHONDRIAL_LARGE_RIBOSOMAL_SUBUNIT
- GO_TRNA_METABOLIC_PROCESS, GO_TRNA_METABOLIC_PROCESS
- GO_TRNA_PROCESSING, GO_TRNA_PROCESSING
- GO_TRNA_WOBBLE_BASE_MODIFICATION, GO_TRNA_WOBBLE_BASE_MODIFICATION
- GO_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT, GO_MITOCHONDRIAL_SMALL_RIBOSOMAL_SUBUNIT
- GO_RNA_MODIFICATION, GO_RNA_MODIFICATION
- GO_TRNA_BINDING, GO_TRNA_BINDING
- GO_RIBOSOMAL_SUBUNIT, GO_RIBOSOMAL_SUBUNIT
- GO_STRUCTURAL_CONSTITUENT_OF_RIBOSOME, GO_STRUCTURAL_CONSTITUENT_OF_RIBOSOME
- GO_TRNA_MODIFICATION, GO_TRNA_MODIFICATION
- GO_DNA_DAMAGE_RESPONSE_DETECTION_OF_DNA_DAMAGE, GO_DNA_DAMAGE_RESPONSE_DETECTION_OF_DNA_DA
- GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY, GO_INTRAMOLECULAR_TRANSFERASE_ACTIVITY
- GO_PSEUDOURIDINE_SYNTHESIS, GO_PSEUDOURIDINE_SYNTHESIS
- GO LARGE RIBOSOMAL SUBUNIT, GO LARGE RIBOSOMAL SUBUNIT
- GO_TRNA_THREONYLCARBAMOYLADENOSINE_METABOLIC_PROCESS, GO_TRNA_THREONYLCARBAMOYLADENOSINE_N
- GO_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING, GO_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING
- GO TRANSLATION REGULATOR ACTIVITY NUCLEIC ACID BINDING, GO TRANSLATION REGULATOR ACTIVITY NUCLEI
- GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_RNA_METABOLIC_PROCESS
- GO_TRNA_3_END_PROCESSING, GO_TRNA_3_END_PROCESSING
- GO_RRNA_METHYLATION, GO_RRNA_METHYLATION
- GO_SMALL_RIBOSOMAL_SUBUNIT, GO_SMALL_RIBOSOMAL_SUBUNIT