## TIONAL\_INITIATION, GO\_TRANSLATIONAL\_INITIATION

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GO_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE, GO_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE
GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_NONSENSE_MEDIATED_DECAY, GO_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_NONSENSE_MEDIATED_DECAY
GO_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM, GO_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
GO_CYTOSOLIC_RIBOSOME, GO_CYTOSOLIC_RIBOSOME
GO_LARGE_RIBOSOMAL_SUBUNIT, GO_LARGE_RIBOSOMAL_SUBUNIT
GO_PROTEIN_TARGETING_TO_MEMBRANE, GO_PROTEIN_TARGETING_TO_MEMBRANE
GO_CYTOSOLIC_LARGE_RIBOSOMAL_SUBUNIT, GO_CYTOSOLIC_LARGE_RIBOSOMAL_SUBUNIT
GO_CYTOPLASMIC_TRANSLATION, GO_CYTOPLASMIC_TRANSLATION
GO_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS, GO_RIBOSOMAL_LARGE_SUBUNIT_BIOGENESIS
GO_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS, GO_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS
GO_TRANSLATION_REGULATOR_ACTIVITY_NUCLEIC_ACID_BINDING, GO_TRANSLATION_REGULATOR_ACTIVITY_NUCLEIC_ACID_BINDING
GO_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING, GO_TRANSLATION_FACTOR_ACTIVITY_RNA_BINDING
GO_TRANSLATION_REGULATOR_ACTIVITY, GO_TRANSLATION_REGULATOR_ACTIVITY
GO_MATURATION_OF_SSU_RRNA, GO_MATURATION_OF_SSU_RRNA
GO_RRNA_BINDING, GO_RRNA_BINDING
GO_SMALL_RIBOSOMAL_SUBUNIT, GO_SMALL_RIBOSOMAL_SUBUNIT
GO_TRANSLATION_INITIATION_FACTOR_ACTIVITY, GO_TRANSLATION_INITIATION_FACTOR_ACTIVITY
GO_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT, GO_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION
GO_POLYSOME, GO_POLYSOME
GO_POLYSOMAL_RIBOSOME, GO_POLYSOMAL_RIBOSOME
GO_POSITIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS, GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS
GO_POSITIVE_REGULATION_OF_TRANSLATION, GO_POSITIVE_REGULATION_OF_TRANSLATION
GO_MATURATION_OF_SSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5_8S_RRNA_LSU_RRNA, GO_MATURATION_OF_SSU_RRNA_FROM_TRICISTRONIC_RRNA_TRANSCRIPT_SSU_RRNA_5
GO_REGULATION_OF_TRANSLATIONAL_INITIATION, GO_REGULATION_OF_TRANSLATIONAL_INITIATION
GO_RIBOSOME_ASSEMBLY, GO_RIBOSOME_ASSEMBLY
GO_TRNA_BINDING, GO_TRNA_BINDING
GO_RIBOSOME_BINDING, GO_RIBOSOME_BINDING
GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GO_PHOSPHATASE_COMPLEX, GO_PHOSPHATASE_COMPLEX
GO_MULTI_ORGANISM_LOCALIZATION, GO_MULTI_ORGANISM_LOCALIZATION
GO_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE, GO_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I
GO_MATURATION_OF_LSU_RRNA, GO_MATURATION_OF_LSU_RRNA
GO_RNA_CAPPING, GO_RNA_CAPPING
GO_RESPONSE_TO_HEAT, GO_RESPONSE_TO_HEAT
GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS
GO_T_CELL_RECEPTOR_SIGNALING_PATHWAY, GO_T_CELL_RECEPTOR_SIGNALING_PATHWAY
GO_POSITIVE_REGULATION_OF_TRANSLATIONAL_INITIATION, GO_POSITIVE_REGULATION_OF_TRANSLATIONAL_INITIATION
GO REGULATION OF VIRAL LIFE CYCLE, GO REGULATION OF VIRAL LIFE CYCLE
GO_CYTOPLASMIC_TRANSLATIONAL_INITIATION, GO_CYTOPLASMIC_TRANSLATIONAL_INITIATION
GO_RESPONSE_TO_UV, GO_RESPONSE_TO_UV
GO_CELLULAR_RESPONSE_TO_RADIATION, GO_CELLULAR_RESPONSE_TO_RADIATION
GO_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GO_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I
GO_TRANSLATION_PREINITIATION_COMPLEX, GO_TRANSLATION_PREINITIATION_COMPLEX
GO_MEMORY, GO_MEMORY
GO_TOR_SIGNALING, GO_TOR_SIGNALING
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GO\_ROUGH\_ENDOPLASMIC\_RETICULUM\_MEMBRANE, GO\_ROUGH\_ENDOPLASMIC\_RETICULUM\_MEMBRANE

GO\_ERROR\_PRONE\_TRANSLESION\_SYNTHESIS, GO\_ERROR\_PRONE\_TRANSLESION\_SYNTHESIS

GO\_RIBOSOMAL\_LARGE\_SUBUNIT\_ASSEMBLY, GO\_RIBOSOMAL\_LARGE\_SUBUNIT\_ASSEMBLY

GO\_VIRAL\_TRANSLATION, GO\_VIRAL\_TRANSLATION

GO\_RIBOSOMAL\_SUBUNIT, GO\_RIBOSOMAL\_SUBUNIT

GO VIRAL GENE EXPRESSION, GO VIRAL GENE EXPRESSION

GO\_STRUCTURAL\_CONSTITUENT\_OF\_RIBOSOME, GO\_STRUCTURAL\_CONSTITUENT\_OF\_RIBOSOME

GO\_CELLULAR\_RESPONSE\_TO\_GLUCOSE\_STARVATION, GO\_CELLULAR\_RESPONSE\_TO\_GLUCOSE\_STARVATION

GO\_PROTEIN\_TAG, GO\_PROTEIN\_TAG

GO\_POSITIVE\_REGULATION\_OF\_APOPTOTIC\_SIGNALING\_PATHWAY, GO\_POSITIVE\_REGULATION\_OF\_APOPTOTIC\_SIGNALING\_PATHWAY

GO\_CELLULAR\_RESPONSE\_TO\_UV, GO\_CELLULAR\_RESPONSE\_TO\_UV

GO\_RRNA\_CONTAINING\_RIBONUCLEOPROTEIN\_COMPLEX\_EXPORT\_FROM\_NUCLEUS, GO\_RRNA\_CONTAINING\_RIBONUCLEOPROTEIN\_COMPLEX\_EXPORT\_FROM\_NUCLEUS

GO\_MODIFICATION\_BY\_SYMBIONT\_OF\_HOST\_MORPHOLOGY\_OR\_PHYSIOLOGY, GO\_MODIFICATION\_BY\_SYMBIONT\_OF\_HOST\_MORPHOLOGY\_OR\_PHYSIOLOGY

GO\_ERROR\_FREE\_TRANSLESION\_SYNTHESIS, GO\_ERROR\_FREE\_TRANSLESION\_SYNTHESIS

GO\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_GAP\_FILLING, GO\_NUCLEOTIDE\_EXCISION\_REPAIR\_DNA\_GAP\_FILLING

GO\_HOST\_CELLULAR\_COMPONENT, GO\_HOST\_CELLULAR\_COMPONENT

GO\_REGULATION\_OF\_UBIQUITIN\_PROTEIN\_TRANSFERASE\_ACTIVITY, GO\_REGULATION\_OF\_UBIQUITIN\_PROTEIN\_TRANSFERASE\_ACTIVITY

GO\_EUKARYOTIC\_TRANSLATION\_INITIATION\_FACTOR\_3\_COMPLEX, GO\_EUKARYOTIC\_TRANSLATION\_INITIATION\_FACTOR\_3\_COMPLEX

GO\_MRNA\_CLEAVAGE, GO\_MRNA\_CLEAVAGE

GO\_RIBOSOMAL\_LARGE\_SUBUNIT\_BINDING, GO\_RIBOSOMAL\_LARGE\_SUBUNIT\_BINDING

GO\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS, GO\_NEGATIVE\_REGULATION\_OF\_RESPONSE\_TO\_ENDOPLASMIC\_RETICULUM\_STRESS

GO\_EUKARYOTIC\_48S\_PREINITIATION\_COMPLEX, GO\_EUKARYOTIC\_48S\_PREINITIATION\_COMPLEX

GO\_RNA\_CAP\_BINDING, GO\_RNA\_CAP\_BINDING

GO\_REGULATION\_OF\_CYTOPLASMIC\_TRANSLATION, GO\_REGULATION\_OF\_CYTOPLASMIC\_TRANSLATION

GO\_MATURATION\_OF\_LSU\_RRNA\_FROM\_TRICISTRONIC\_RRNA\_TRANSCRIPT\_SSU\_RRNA\_5\_8S\_RRNA\_LSU\_RRNA, GO\_MATURATION\_OF\_LSU\_RRNA\_FROM\_TRICISTRONIC\_RRNA\_TRANSCRIPT\_SSU\_RRNA\_5

GO\_FORMATION\_OF\_CYTOPLASMIC\_TRANSLATION\_INITIATION\_COMPLEX, GO\_FORMATION\_OF\_CYTOPLASMIC\_TRANSLATION\_INITIATION\_COMPLEX

GO\_NEGATIVE\_REGULATION\_OF\_CELL\_PROJECTION\_ORGANIZATION, GO\_NEGATIVE\_REGULATION\_OF\_CELL\_PROJECTION\_ORGANIZATION

GO\_IRES\_DEPENDENT\_VIRAL\_TRANSLATIONAL\_INITIATION, GO\_IRES\_DEPENDENT\_VIRAL\_TRANSLATIONAL\_INITIATION

GO\_REGULATION\_OF\_DEOXYRIBONUCLEASE\_ACTIVITY, GO\_REGULATION\_OF\_DEOXYRIBONUCLEASE\_ACTIVITY