

GENE_EXPRESSION, GO_VIRAL_GENE_EXPRESSION

- GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
- GO_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM, GO_PROTEIN_LOCALIZATION_TO_ENDOPLASMIC_RETICULUM
- GO_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE, GO_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE
- GO_PROTEIN_TARGETING_TO_MEMBRANE, GO_PROTEIN_TARGETING_TO_MEMBRANE
- GO_NCRNA_TRANSCRIPTION, GO_NCRNA_TRANSCRIPTION
- GO_SNRNA_TRANSCRIPTION, GO_SNRNA_TRANSCRIPTION
- GO_TRANSLATIONAL_INITIATION, GO_TRANSLATIONAL_INITIATION
- GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
- GO_RNA_POLYMERASE_II_HOLOENZYME, GO_RNA_POLYMERASE_II_HOLOENZYME
- GO_CYTOSOLIC_LARGE_RIBOSOMAL_SUBUNIT, GO_CYTOSOLIC_LARGE_RIBOSOMAL_SUBUNIT
- GO_CYTOSOLIC_RIBOSOME, GO_CYTOSOLIC_RIBOSOME
- GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS, GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS
- GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
- GO_PROTEIN_IMPORT, GO_PROTEIN_IMPORT
- GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING, GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING
- GO_RNA_POLYMERASE_ACTIVITY, GO_RNA_POLYMERASE_ACTIVITY
- GO_RNA_CAPPING, GO_RNA_CAPPING
- GO_TYPE_I_INTERFERON_PRODUCTION, GO_TYPE_I_INTERFERON_PRODUCTION
- GO_CYTOPLASMIC_TRANSLATION, GO_CYTOPLASMIC_TRANSLATION
- GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX, GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX
- GO_P_BODY, GO_P_BODY
- GO_NEGATIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_REGULATION_OF_MRNA_3_END_PROCESSING, GO_REGULATION_OF_MRNA_3_END_PROCESSING
- GO_TRANSCRIPTION_BY_RNA_POLYMERASE_III, GO_TRANSCRIPTION_BY_RNA_POLYMERASE_III
- GO_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GO_TRANSCRIPTION_BY_RNA_POLYMERASE_I
- GO_THYROID_HORMONE_RECEPTOR_BINDING, GO_THYROID_HORMONE_RECEPTOR_BINDING
- GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION, GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION
- GO_TRANSLATION_INITIATION_FACTOR_BINDING, GO_TRANSLATION_INITIATION_FACTOR_BINDING
- GO_REGULATION_OF_VIRAL_TRANSCRIPTION, GO_REGULATION_OF_VIRAL_TRANSCRIPTION
- GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
- GO_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT, GO_CYTOSOLIC_SMALL_RIBOSOMAL_SUBUNIT
- GO_NEGATIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS
- GO_POLYSOME, GO_POLYSOME
- GO_POSITIVE_REGULATION_OF_AUTOPHAGY, GO_POSITIVE_REGULATION_OF_AUTOPHAGY
- GO_MOVEMENT_IN_ENVIRONMENT_OF_OTHER_ORGANISM_INVOLVED_IN_SYMBIOTIC_INTERACTION, GO_MOVEMENT_IN_ENVIRONMENT_OF_OTHER_ORGANISM_INVOLVED_IN_SYMBIOTIC_INTERACTION
- GO_TERMINATION_OF_RNA_POLYMERASE_I_TRANSCRIPTION, GO_TERMINATION_OF_RNA_POLYMERASE_I_TRANSCRIPTION
- GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_I_PROMOTER, GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_I_PROMOTER
- GO_REGULATION_OF_TRANSLATIONAL_INITIATION, GO_REGULATION_OF_TRANSLATIONAL_INITIATION
- GO_POLYSOMAL_RIBOSOME, GO_POLYSOMAL_RIBOSOME
- GO_REGULATION_OF_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, GO_REGULATION_OF_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
- GO_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS, GO_RIBOSOMAL_SMALL_SUBUNIT_BIOGENESIS
- GO_REGULATION_OF_MRNA_POLYADENYLATION, GO_REGULATION_OF_MRNA_POLYADENYLATION
- GO_TRANSCRIPTION_COACTIVATOR_BINDING, GO_TRANSCRIPTION_COACTIVATOR_BINDING
- GO_CARBOXY_TERMINAL_DOMAIN_PROTEIN_KINASE_COMPLEX, GO_CARBOXY_TERMINAL_DOMAIN_PROTEIN_KINASE_COMPLEX
- GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
- GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I_PROMOTER, GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_I_PROMOTER
- GO_REGULATION_OF_NUCLEOBASE_CONTAINING_COMPOUND_TRANSPORT, GO_REGULATION_OF_NUCLEOBASE_CONTAINING_COMPOUND_TRANSPORT
- GO_VIRAL_TRANSLATION, GO_VIRAL_TRANSLATION
- GO_TRANSCRIPTION_COFACTOR_BINDING, GO_TRANSCRIPTION_COFACTOR_BINDING
- GO_SIGNAL_SEQUENCE_BINDING, GO_SIGNAL_SEQUENCE_BINDING
- GO_NLS_BEARING_PROTEIN_IMPORT_INTO_NUCLEUS, GO_NLS_BEARING_PROTEIN_IMPORT_INTO_NUCLEUS
- GO_IRES_DEPENDENT_VIRAL_TRANSLATIONAL_INITIATION, GO_IRES_DEPENDENT_VIRAL_TRANSLATIONAL_INITIATION
- GO_TRANSCRIPTION_FACTOR_TFIID_COMPLEX, GO_TRANSCRIPTION_FACTOR_TFIID_COMPLEX
- GO_VITAMIN_D_RECEPTOR_BINDING, GO_VITAMIN_D_RECEPTOR_BINDING
- GO_RNA_POLYMERASE_II_ACTIVITY, GO_RNA_POLYMERASE_II_ACTIVITY
- GO_TRANSLATION_INITIATION_FACTOR_ACTIVITY, GO_TRANSLATION_INITIATION_FACTOR_ACTIVITY
- GO_POSITIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY, GO_POSITIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY
- GO_SEQUENCE_SPECIFIC_MRNA_BINDING, GO_SEQUENCE_SPECIFIC_MRNA_BINDING
- GO_VIRAL_RNA_GENOME_REPLICATION, GO_VIRAL_RNA_GENOME_REPLICATION
- GO_NUCLEAR_EXPORT_SIGNAL_RECEPTOR_ACTIVITY, GO_NUCLEAR_EXPORT_SIGNAL_RECEPTOR_ACTIVITY
- GO_RRNA_CONTAINING_RIBONUCLEOPROTEIN_COMPLEX_EXPORT_FROM_NUCLEUS, GO_RRNA_CONTAINING_RIBONUCLEOPROTEIN_COMPLEX_EXPORT_FROM_NUCLEUS
- GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS_BY_HOST