

EAR_EXPORT, GO_NUCLEAR_EXPORT

GO_ESTABLISHMENT_OF_RNA_LOCALIZATION, GO_ESTABLISHMENT_OF_RNA_LOCALIZATION
GO_RNA_EXPORT_FROM_NUCLEUS, GO_RNA_EXPORT_FROM_NUCLEUS
GO_SPLICEOSOMAL_COMPLEX, GO_SPLICEOSOMAL_COMPLEX
GO_MRNA_EXPORT_FROM_NUCLEUS, GO_MRNA_EXPORT_FROM_NUCLEUS
GO_MRNA_TRANSPORT, GO_MRNA_TRANSPORT
GO_REGULATION_OF_GENE_SILENCING, GO_REGULATION_OF_GENE_SILENCING
GO_REGULATION_OF_MRNA_PROCESSING, GO_REGULATION_OF_MRNA_PROCESSING
GO_RNA_3_END_PROCESSING, GO_RNA_3_END_PROCESSING
GO_KINETOCHORE, GO_KINETOCHORE
GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
GO_CATALYTIC_STEP_2_SPLICEOSOME, GO_CATALYTIC_STEP_2_SPLICEOSOME
GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
GO_CHROMOSOME_TELOMERIC_REGION, GO_CHROMOSOME_TELOMERIC_REGION
GO_REGULATION_OF_MRNA_SPLICING_VIA_SPLICEOSOME, GO_REGULATION_OF_MRNA_SPLICING_VIA_SPLICEOSOME
GO_REGULATION_OF_RNA_SPLICING, GO_REGULATION_OF_RNA_SPLICING
GO_U2_TYPE_SPLICEOSOMAL_COMPLEX, GO_U2_TYPE_SPLICEOSOMAL_COMPLEX
GO_MRNA_3_END_PROCESSING, GO_MRNA_3_END_PROCESSING
GO_REGULATION_OF_TELOMERE_MAINTENANCE, GO_REGULATION_OF_TELOMERE_MAINTENANCE
GO_NUCLEAR_PORE, GO_NUCLEAR_PORE
GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION, GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION
GO_NCRNA_EXPORT_FROM_NUCLEUS, GO_NCRNA_EXPORT_FROM_NUCLEUS
GO_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME, GO_ALTERNATIVE_MRNA_SPLICING_VIA_SPLICEOSOME
GO_CONDENSED_CHROMOSOME_CENTROMERIC_REGION, GO_CONDENSED_CHROMOSOME_CENTROMERIC_REGION
GO_DNA_TEMPLATED_TRANSCRIPTION_TERMINATION, GO_DNA_TEMPLATED_TRANSCRIPTION_TERMINATION
GO_MULTI_ORGANISM_LOCALIZATION, GO_MULTI_ORGANISM_LOCALIZATION
GO_PROTEIN_SUMOYLATION, GO_PROTEIN_SUMOYLATION
GO_SPLICEOSOMAL_COMPLEX_ASSEMBLY, GO_SPLICEOSOMAL_COMPLEX_ASSEMBLY
GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GO_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS, GO_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
GO_REGULATION_OF_TRANSLATIONAL_INITIATION, GO_REGULATION_OF_TRANSLATIONAL_INITIATION
GO_MITOTIC_SPINDLE_ORGANIZATION, GO_MITOTIC_SPINDLE_ORGANIZATION
GO_NEGATIVE_REGULATION_OF_MRNA_SPLICING_VIA_SPLICEOSOME, GO_NEGATIVE_REGULATION_OF_MRNA_SPLICING_VIA_SPLICEOSOME
GO_HOST_CELLULAR_COMPONENT, GO_HOST_CELLULAR_COMPONENT
GO_CELLULAR_RESPONSE_TO_HEAT, GO_CELLULAR_RESPONSE_TO_HEAT
GO_RESPONSE_TO_HEAT, GO_RESPONSE_TO_HEAT
GO_PROTEIN_IMPORT, GO_PROTEIN_IMPORT
GO_NUCLEAR_PERIPHERY, GO_NUCLEAR_PERIPHERY
GO_TERMINATION_OF_RNA_POLYMERASE_II_TRANSCRIPTION, GO_TERMINATION_OF_RNA_POLYMERASE_II_TRANSCRIPTION
GO_MRNA_SPLICE_SITE_SELECTION, GO_MRNA_SPLICE_SITE_SELECTION
GO_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE, GO_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE
GO_NEGATIVE_REGULATION_OF_RNA_SPLICING, GO_NEGATIVE_REGULATION_OF_RNA_SPLICING
GO_RNA_HELICASE_ACTIVITY, GO_RNA_HELICASE_ACTIVITY
GO_NUCLEUS_ORGANIZATION, GO_NUCLEUS_ORGANIZATION
GO_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS, GO_REGULATION_OF_CARBOHYDRATE_METABOLIC_PROCESS
GO_NEGATIVE_REGULATION_OF_MRNA_PROCESSING, GO_NEGATIVE_REGULATION_OF_MRNA_PROCESSING
GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
GO_IMPORT_INTO_NUCLEUS, GO_IMPORT_INTO_NUCLEUS
GO_NEGATIVE_REGULATION_OF_MRNA_METABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_MRNA_METABOLIC_PROCESS
GO_PML_BODY, GO_PML_BODY
GO_NUCLEAR_MATRIX, GO_NUCLEAR_MATRIX
GO_NUCLEOCYTOPLASMIC_CARRIER_ACTIVITY, GO_NUCLEOCYTOPLASMIC_CARRIER_ACTIVITY
GO_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY, GO_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY
GO_MRNA_CIS_SPLICING_VIA_SPLICEOSOME, GO_MRNA_CIS_SPLICING_VIA_SPLICEOSOME
GO_CAJAL_BODY, GO_CAJAL_BODY
GO_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GO_NUCLEAR_PORE_NUCLEAR_BASKET, GO_NUCLEAR_PORE_NUCLEAR_BASKET
GO_PROTEIN_STABILIZATION, GO_PROTEIN_STABILIZATION
GO_REGULATION_OF_HISTONE_MODIFICATION, GO_REGULATION_OF_HISTONE_MODIFICATION
GO_SINGLE_STRANDED_RNA_BINDING, GO_SINGLE_STRANDED_RNA_BINDING
GO_TELOMERIC_DNA_BINDING, GO_TELOMERIC_DNA_BINDING
GO_NUCLEAR_PORE_ORGANIZATION, GO_NUCLEAR_PORE_ORGANIZATION
GO_HISTONE_MRNA_METABOLIC_PROCESS, GO_HISTONE_MRNA_METABOLIC_PROCESS
GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION
GO_RRNA_CONTAINING_RIBONUCLEOPROTEIN_COMPLEX_EXPORT_FROM_NUCLEUS, GO_RRNA_CONTAINING_RIBONUCLEOPROTEIN_COMPLEX_EXPORT_FROM_NUCLEUS
GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING, GO_NUCLEAR_LOCALIZATION_SEQUENCE_BINDING
GO_CHROMOSOME_LOCALIZATION, GO_CHROMOSOME_LOCALIZATION
GO_RAN_GTPASE_BINDING, GO_RAN_GTPASE_BINDING
GO_TRANSLATION_INITIATION_FACTOR_BINDING, GO_TRANSLATION_INITIATION_FACTOR_BINDING
GO_NUCLEAR_ENVELOPE_ORGANIZATION, GO_NUCLEAR_ENVELOPE_ORGANIZATION
GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT, GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT
GO_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR, GO_PRERIBOSOME_LARGE_SUBUNIT_PRECURSOR
GO_RNA_CAPPING, GO_RNA_CAPPING
GO_REGULATION_OF_ATP_METABOLIC_PROCESS, GO_REGULATION_OF_ATP_METABOLIC_PROCESS
GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT, GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT
GO_REGULATORY_RNA_BINDING, GO_REGULATORY_RNA_BINDING
GO_CATALYTIC_STEP_1_SPLICEOSOME, GO_CATALYTIC_STEP_1_SPLICEOSOME
GO_PRE_MRNA_BINDING, GO_PRE_MRNA_BINDING
GO_PROTEIN_HETEROTETRAMERIZATION, GO_PROTEIN_HETEROTETRAMERIZATION
GO_NUCLEAR_INCLUSION_BODY, GO_NUCLEAR_INCLUSION_BODY
GO_TRANSCRIPTION_EXPORT_COMPLEX, GO_TRANSCRIPTION_EXPORT_COMPLEX
GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS, GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS
GO_REGULATION_OF_CELL_CYCLE_ARREST, GO_REGULATION_OF_CELL_CYCLE_ARREST
GO_RNAI_EFFECTOR_COMPLEX, GO_RNAI_EFFECTOR_COMPLEX
GO_CYTOPLASMIC_STRESS_GRANULE, GO_CYTOPLASMIC_STRESS_GRANULE
GO_REGULATION_OF_SPINDLE_ORGANIZATION, GO_REGULATION_OF_SPINDLE_ORGANIZATION
GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_ELONGATION
GO_G1_DNA_DAMAGE_CHECKPOINT, GO_G1_DNA_DAMAGE_CHECKPOINT
GO_HYALURONAN_CATABOLIC_PROCESS, GO_HYALURONAN_CATABOLIC_PROCESS
GO_PROTEIN_N_TERMINUS_BINDING, GO_PROTEIN_N_TERMINUS_BINDING
GO_POSITIVE_REGULATION_OF_RNA_SPLICING, GO_POSITIVE_REGULATION_OF_RNA_SPLICING
GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION, GO_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_LOCALIZATION
GO_MRNA_3_UTR_BINDING, GO_MRNA_3_UTR_BINDING
GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
GO_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS, GO_POLY_A_PLUS_MRNA_EXPORT_FROM_NUCLEUS
GO_LYMPH_VESSEL_MORPHOGENESIS, GO_LYMPH_VESSEL_MORPHOGENESIS
GO_ANDROGEN_RECEPTOR_BINDING, GO_ANDROGEN_RECEPTOR_BINDING
GO_REGULATION_OF_HISTONE_PHOSPHORYLATION, GO_REGULATION_OF_HISTONE_PHOSPHORYLATION
GO_P53_BINDING, GO_P53_BINDING
GO_REGULATION_OF_TELOMERASE_ACTIVITY, GO_REGULATION_OF_TELOMERASE_ACTIVITY
GO_CARBOHYDRATE_CATABOLIC_PROCESS, GO_CARBOHYDRATE_CATABOLIC_PROCESS
GO_REGULATION_OF_MRNA_3_END_PROCESSING, GO_REGULATION_OF_MRNA_3_END_PROCESSING
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST
GO_REGULATION_OF_CENTRIOLE_REPLICATION, GO_REGULATION_OF_CENTRIOLE_REPLICATION
GO_EXON_EXON_JUNCTION_COMPLEX, GO_EXON_EXON_JUNCTION_COMPLEX
GO_PROTEIN_HETEROOLIGOMERIZATION, GO_PROTEIN_HETEROOLIGOMERIZATION
GO_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR, GO_PRERIBOSOME_SMALL_SUBUNIT_PRECURSOR
GO_MAINTENANCE_OF_PROTEIN_LOCATION, GO_MAINTENANCE_OF_PROTEIN_LOCATION
GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE, GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE
GO_MAINTENANCE_OF_PROTEIN_LOCATION_IN_CELL, GO_MAINTENANCE_OF_PROTEIN_LOCATION_IN_CELL
GO_POSITIVE_REGULATION_OF_HISTONE_DEACETYLATION, GO_POSITIVE_REGULATION_OF_HISTONE_DEACETYLATION
GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION, GO_RNA_SPLICING_VIA_ENDONUCLEOLYTIC_CLEAVAGE_AND_LIGATION
GO_HISTONE_PHOSPHORYLATION, GO_HISTONE_PHOSPHORYLATION
GO_P_BODY, GO_P_BODY
GO_RIBOSOMAL_LARGE_SUBUNIT_EXPORT_FROM_NUCLEUS, GO_RIBOSOMAL_LARGE_SUBUNIT_EXPORT_FROM_NUCLEUS
GO_ER_NUCLEUS_SIGNALING_PATHWAY, GO_ER_NUCLEUS_SIGNALING_PATHWAY