

BRANE_ORGANIZATION, GO_MITOCHONDRIAL_MEMBRANE_ORGANIZATION

GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
GO_OXIDATIVE_PHOSPHORYLATION, GO_OXIDATIVE_PHOSPHORYLATION
GO_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GO_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
GO_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT, GO_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT
GO_CELLULAR_RESPIRATION, GO_CELLULAR_RESPIRATION
GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE
GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE
GO_ORGANELLE_FUSION, GO_ORGANELLE_FUSION
GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT, GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT
GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
GO_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_REGULATION_OF_MITOCHONDRION_ORGANIZATION
GO_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN, GO_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN
GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX, GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX
GO_PURINE_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GO_PURINE_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
GO_ELECTRON_TRANSPORT_CHAIN, GO_ELECTRON_TRANSPORT_CHAIN
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
GO_B_CELL_HOMEOSTASIS, GO_B_CELL_HOMEOSTASIS
GO_PROTEIN_TARGETING_TO_MITOCHONDRION, GO_PROTEIN_TARGETING_TO_MITOCHONDRION
GO_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GO_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
GO_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GO_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
GO_ORGANELLE_DISASSEMBLY, GO_ORGANELLE_DISASSEMBLY
GO_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
GO_CRISTAE_FORMATION, GO_CRISTAE_FORMATION
GO_ATP_BIOSYNTHETIC_PROCESS, GO_ATP_BIOSYNTHETIC_PROCESS
GO_POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
GO_APOPTOTIC_MITOCHONDRIAL_CHANGES, GO_APOPTOTIC_MITOCHONDRIAL_CHANGES
GO_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GO_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
GO_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
GO_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR, GO_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR
GO_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE, GO_IRE1_MEDIATED_UNFOLDED_PROTEIN_RESPONSE
GO_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS, GO_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS
GO_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GO_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX, GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX
GO_CHAPERONE_BINDING, GO_CHAPERONE_BINDING
GO_POSITIVE_REGULATION_OF_MEMBRANE_PERMEABILITY, GO_POSITIVE_REGULATION_OF_MEMBRANE_PERMEABILITY
GO_PEPTIDYL_THREONINE_MODIFICATION, GO_PEPTIDYL_THREONINE_MODIFICATION
GO_TAU_PROTEIN_KINASE_ACTIVITY, GO_TAU_PROTEIN_KINASE_ACTIVITY
GO_HYDROGEN_PEROXIDE_BIOSYNTHETIC_PROCESS, GO_HYDROGEN_PEROXIDE_BIOSYNTHETIC_PROCESS
GO_AXON_CYTOPLASM, GO_AXON_CYTOPLASM
GO_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION, GO_CELLULAR_RESPONSE_TO_GLUCOSE_STARVATION
GO_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE, GO_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE
GO_AXO_DENDRITIC_TRANSPORT, GO_AXO_DENDRITIC_TRANSPORT
GO_REGULATION_OF_NEUROTRANSMITTER_SECRETION, GO_REGULATION_OF_NEUROTRANSMITTER_SECRETION
GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_OUTER_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_OUTER_MEMBRANE
GO_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT, GO_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN, GO_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_PROTON_TRANSPORTING_DOMAIN
GO_POSITIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY
GO_REGULATION_OF_SYNAPTIC_TRANSMISSION_DOPAMINERGIC, GO_REGULATION_OF_SYNAPTIC_TRANSMISSION_DOPAMINERGIC
GO_REGULATION_OF_MEMBRANE_PERMEABILITY, GO_REGULATION_OF_MEMBRANE_PERMEABILITY
GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE, GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE
GO_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION, GO_MITOCHONDRIAL_OUTER_MEMBRANE_PERMEABILIZATION
GO_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GO_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
GO_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
GO_PIGMENT_GRANULE, GO_PIGMENT_GRANULE
GO_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE, GO_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE
GO_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY_INVOLVED_IN_APOPTOTIC_PROCESS, GO_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_PERMEABILITY_INVOLVED_IN_APOPTOTIC_PROCESS
GO_MYELOID_CELL_HOMEOSTASIS, GO_MYELOID_CELL_HOMEOSTASIS
GO_RIPOPTOSOME, GO_RIPOPTOSOME
GO_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM, GO_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM
GO_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DIFFERENTIATION, GO_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DIFFERENTIATION
GO_POSITIVE_REGULATION_OF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_CYSINE_TYPE_ENDOPEPTIDASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL, GO_POSITIVE_REGULATION_OF_MEMBRANE_POTENTIAL
GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE, GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_MEMBRANE
GO_FEMALE_GENITALIA_DEVELOPMENT, GO_FEMALE_GENITALIA_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GO_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
GO_AUTOPHAGY_OF_MITOCHONDRION, GO_AUTOPHAGY_OF_MITOCHONDRION
GO_SYNAPTIC_VESICLE_ENDOCYTOSIS, GO_SYNAPTIC_VESICLE_ENDOCYTOSIS
GO_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT, GO_COLUMNAR_CUBOIDAL_EPITHELIAL_CELL_DEVELOPMENT
GO_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE, GO_ENDOPLASMIC_RETICULUM_UNFOLDED_PROTEIN_RESPONSE
GO_SUBSTANTIA_NIGRA_DEVELOPMENT, GO_SUBSTANTIA_NIGRA_DEVELOPMENT
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRIAL_MEMBRANE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRIAL_MEMBRANE
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
GO_PROTON_CHANNEL_ACTIVITY, GO_PROTON_CHANNEL_ACTIVITY
GO_PROTEIN_TRANSMEMBRANE_IMPORT_INTO_INTRACELLULAR_ORGANELLE, GO_PROTEIN_TRANSMEMBRANE_IMPORT_INTO_INTRACELLULAR_ORGANELLE
GO_ENDOPLASMIC_RETICULUM_CALCIIUM_ION_HOMEOSTASIS, GO_ENDOPLASMIC_RETICULUM_CALCIIUM_ION_HOMEOSTASIS