

ULAR RESPIRATION, GOBP_CELLULAR RESPIRATION

- GOBP_OXIDATIVE_PHOSPHORYLATION, GOBP_OXIDATIVE_PHOSPHORYLATION
- GOCC_INNER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX, GOCC_INNER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX
- GOBP_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN, GOBP_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN
- GOBP_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT, GOBP_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT
- GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY, GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY
- GOBP_ELECTRON_TRANSPORT_CHAIN, GOBP_ELECTRON_TRANSPORT_CHAIN
- GOCC_RESPIRASOME, GOCC_RESPIRASOME
- GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR
- GOMF_NAD_P_H_DEHYDROGENASE_QUINONE_ACTIVITY, GOMF_NAD_P_H_DEHYDROGENASE_QUINONE_ACTIVITY
- GOCC_RESPIRATORY_CHAIN_COMPLEX, GOCC_RESPIRATORY_CHAIN_COMPLEX
- GOCC_NADH_DEHYDROGENASE_COMPLEX, GOCC_NADH_DEHYDROGENASE_COMPLEX
- GOBP_AEROBIC_RESPIRATION, GOBP_AEROBIC_RESPIRATION
- GOMF_ELECTRON_TRANSFER_ACTIVITY, GOMF_ELECTRON_TRANSFER_ACTIVITY
- GOCC_OXIDOREDUCTASE_COMPLEX, GOCC_OXIDOREDUCTASE_COMPLEX
- GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE
- GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE
- GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE
- GOCC_ORGANELLE_ENVELOPE_LUMEN, GOCC_ORGANELLE_ENVELOPE_LUMEN
- GOBP_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GOBP_ORGANELLE_DISASSEMBLY, GOBP_ORGANELLE_DISASSEMBLY
- GOBP_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
- GOBP_POSITIVE_REGULATION_OF_AUTOPHAGY, GOBP_POSITIVE_REGULATION_OF_AUTOPHAGY
- GOBP_TRICARBOXYLIC_ACID_CYCLE, GOBP_TRICARBOXYLIC_ACID_CYCLE
- GOBP_PEROXISOMAL_TRANSPORT, GOBP_PEROXISOMAL_TRANSPORT
- GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GOBP_PROTEIN_TARGETING_TO_MITOCHONDRION, GOBP_PROTEIN_TARGETING_TO_MITOCHONDRION
- GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_CH_GROUP_OF_DONORS
- GOBP_HYDROGEN_PEROXIDE_CATABOLIC_PROCESS, GOBP_HYDROGEN_PEROXIDE_CATABOLIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
- GOMF_ORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_ORGANIC_ANION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
- GOBP_NUCLEOSIDE_BISPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOSIDE_BISPHOSPHATE_BIOSYNTHETIC_PROCESS
- GOMF_QUINONE_BINDING, GOMF_QUINONE_BINDING
- GOBP_REGULATION_OF_NMDA_RECEPTOR_ACTIVITY, GOBP_REGULATION_OF_NMDA_RECEPTOR_ACTIVITY
- GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION, GOBP_REGULATION_OF_MITOCHONDRIAL_FISSION
- GOMF_C4_DICARBOXYLATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_C4_DICARBOXYLATE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
- GOBP_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_DEHYDROGENASE, GOBP_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_DEHYDROGENASE
- GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION, GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION
- GOBP_HYPEROSMOTIC_SALINITY_RESPONSE, GOBP_HYPEROSMOTIC_SALINITY_RESPONSE
- GOCC_TRICARBOXYLIC_ACID_CYCLE_ENZYME_COMPLEX, GOCC_TRICARBOXYLIC_ACID_CYCLE_ENZYME_COMPLEX
- GOBP_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_ASSEMBLY, GOBP_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX_ASSEMBLY
- GOBP_NEGATIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS
- GOMF_L_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_L_AMINO_ACID_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
- GOBP_AMYLOID_BETA_CLEARANCE, GOBP_AMYLOID_BETA_CLEARANCE
- GOBP_MITOCHONDRIAL_FISSION, GOBP_MITOCHONDRIAL_FISSION