

_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_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY, GOBP_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY
- GOBP_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT, GOBP_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT
- GOBP_CELLULAR_RESPIRATION, GOBP_CELLULAR_RESPIRATION
- GOBP_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GOCC_ORGANELLE_ENVELOPE_LUMEN, GOCC_ORGANELLE_ENVELOPE_LUMEN
- GOBP_ELECTRON_TRANSPORT_CHAIN, GOBP_ELECTRON_TRANSPORT_CHAIN
- GOCC_RESPIRASOME, GOCC_RESPIRASOME
- GOBP_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE
- GOCC_RESPIRATORY_CHAIN_COMPLEX, GOCC_RESPIRATORY_CHAIN_COMPLEX
- GOCC_NADH_DEHYDROGENASE_COMPLEX, GOCC_NADH_DEHYDROGENASE_COMPLEX
- GOCC_OXIDOREDUCTASE_COMPLEX, GOCC_OXIDOREDUCTASE_COMPLEX
- GOMF_NAD_P_H_DEHYDROGENASE_QUINONE_ACTIVITY, GOMF_NAD_P_H_DEHYDROGENASE_QUINONE_ACTIVITY
- GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY, GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY
- GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT, GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
- GOMF_ELECTRON_TRANSFER_ACTIVITY, GOMF_ELECTRON_TRANSFER_ACTIVITY
- GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- 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
- GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GOBP_CYTOCHROME_COMPLEX_ASSEMBLY, GOBP_CYTOCHROME_COMPLEX_ASSEMBLY
- GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE
- GOBP_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GOBP_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_APOPTOTIC_MITOCHONDRIAL_CHANGES, GOBP_APOPTOTIC_MITOCHONDRIAL_CHANGES
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H
- GOBP_CRISTAE_FORMATION, GOBP_CRISTAE_FORMATION
- GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX, GOCC_PROTON_TRANSPORTING_TWO_SECTOR_ATPASE_COMPLEX
- GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_INNER_MEMBRANE
- GOCC_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX, GOCC_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX
- GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
- GOBP_AEROBIC_RESPIRATION, GOBP_AEROBIC_RESPIRATION
- GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOBP_RESPIRATORY_CHAIN_COMPLEX_IV_ASSEMBLY, GOBP_RESPIRATORY_CHAIN_COMPLEX_IV_ASSEMBLY
- GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_REGULATION_OF_CELL_CYCLE_ARREST, GOBP_REGULATION_OF_CELL_CYCLE_ARREST
- GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOBP_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS, GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_UBIQUINOL_TO_CYTOCHROME_C, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_UBIQUINOL_TO_CYTOCHROME_C
- GOBP_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE, GOBP_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE
- GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GOBP_UBIQUINONE_METABOLIC_PROCESS, GOBP_UBIQUINONE_METABOLIC_PROCESS
- GOMF_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM, GOMF_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM
- GOMF_NAD_BINDING, GOMF_NAD_BINDING
- GOMF_PRIMARY_ACTIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GOMF_PRIMARY_ACTIVE_TRANSMEMBRANE_TRANSPORTER_ACTIVITY