

PHOSPHORYLATION, GO_OXIDATIVE_PHOSPHORYLATION

- GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY, GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY
- GO_INNER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX, GO_INNER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX
- GO_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT, GO_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT
- GO_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN, GO_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN
- GO_CELLULAR_RESPIRATION, GO_CELLULAR_RESPIRATION
- GO_RESPIRASOME, GO_RESPIRASOME
- GO_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY, GO_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY
- GO_RESPIRATORY_CHAIN_COMPLEX, GO_RESPIRATORY_CHAIN_COMPLEX
- GO_AEROBIC_RESPIRATION, GO_AEROBIC_RESPIRATION
- GO_NADH_DEHYDROGENASE_COMPLEX, GO_NADH_DEHYDROGENASE_COMPLEX
- GO_ELECTRON_TRANSPORT_CHAIN, GO_ELECTRON_TRANSPORT_CHAIN
- GO_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE, GO_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE
- GO_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GO_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GO_NADH_DEHYDROGENASE_ACTIVITY, GO_NADH_DEHYDROGENASE_ACTIVITY
- GO_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GO_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GO_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GO_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE, GO_INTRINSIC_COMPONENT_OF_MITOCHONDRIAL_MEMBRANE
- GO_PURINE_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GO_PURINE_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GO_CYTOCHROME_COMPLEX_ASSEMBLY, GO_CYTOCHROME_COMPLEX_ASSEMBLY
- GO_OXIDOREDUCTASE_COMPLEX, GO_OXIDOREDUCTASE_COMPLEX
- GO_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GO_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GO_PROTEIN_TARGETING_TO_MITOCHONDRION, GO_PROTEIN_TARGETING_TO_MITOCHONDRION
- GO_RESPIRATORY_CHAIN_COMPLEX_IV_ASSEMBLY, GO_RESPIRATORY_CHAIN_COMPLEX_IV_ASSEMBLY
- GO_CRISTAE_FORMATION, GO_CRISTAE_FORMATION
- GO_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT, GO_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
- GO_ATP_BIOSYNTHETIC_PROCESS, GO_ATP_BIOSYNTHETIC_PROCESS
- GO_ORGANELLE_ENVELOPE_LUMEN, GO_ORGANELLE_ENVELOPE_LUMEN
- GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H_QUINONE_OR_SIMILAR_COMPOUND_AS_ACCEPTOR
- GO_POSITIVE_REGULATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY
- GO_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR, GO_PHOSPHOTRANSFERASE_ACTIVITY_PHOSPHATE_GROUP_AS_ACCEPTOR
- GO_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS, GO_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_ADP_METABOLIC_PROCESS, GO_ADP_METABOLIC_PROCESS
- GO_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_PEPTIDASE_ACTIVITY
- GO_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GO_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GO_NADH_METABOLIC_PROCESS, GO_NADH_METABOLIC_PROCESS
- GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX, GO_PROTON_TRANSPORTING_ATP_SYNTHASE_COMPLEX
- GO_NUCLEOTIDE_PHOSPHORYLATION, GO_NUCLEOTIDE_PHOSPHORYLATION
- GO_REGULATION_OF_ATP_METABOLIC_PROCESS, GO_REGULATION_OF_ATP_METABOLIC_PROCESS
- GO_TRICARBOXYLIC_ACID_CYCLE, GO_TRICARBOXYLIC_ACID_CYCLE
- GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_NAD_P_H
- GO_PYRUVATE_METABOLIC_PROCESS, GO_PYRUVATE_METABOLIC_PROCESS
- GO_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS, GO_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS
- GO_RESPIRATORY_CHAIN_COMPLEX_III, GO_RESPIRATORY_CHAIN_COMPLEX_III
- GO_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY, GO_REGULATION_OF_GENERATION_OF_PRECURSOR_METABOLITES_AND_ENERGY
- GO_ELECTRON_TRANSFER_ACTIVITY, GO_ELECTRON_TRANSFER_ACTIVITY
- GO_NUCLEOBASE_CONTAINING_COMPOUND_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_NUCLEOBASE_CONTAINING_COMPOUND_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
- GO_ORGANOPHOSPHATE_ESTER_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_ORGANOPHOSPHATE_ESTER_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
- GO_AEROBIC_ELECTRON_TRANSPORT_CHAIN, GO_AEROBIC_ELECTRON_TRANSPORT_CHAIN
- GO_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM, GO_PROTON_TRANSPORTING_ATP_SYNTHASE_ACTIVITY_ROTATIONAL_MECHANISM
- GO_DOPAMINE_METABOLIC_PROCESS, GO_DOPAMINE_METABOLIC_PROCESS
- GO_POSITIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY
- GO_IRON_SULFUR_CLUSTER_ASSEMBLY, GO_IRON_SULFUR_CLUSTER_ASSEMBLY