

_PHOSPHORYLATION, GOBP_OXIDATIVE_PHOSPHORYLATION

- GOBP_CELLULAR_RESPIRATION, GOBP_CELLULAR_RESPIRATION
- GOBP_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT, GOBP_ATP_SYNTHESIS_COUPLED_ELECTRON_TRANSPORT
- GOBP_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN, GOBP_RESPIRATORY_ELECTRON_TRANSPORT_CHAIN
- GOBP_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY, GOBP_NADH_DEHYDROGENASE_COMPLEX_ASSEMBLY
- GOBP_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION, GOBP_INNER_MITOCHONDRIAL_MEMBRANE_ORGANIZATION
- GOBP_AEROBIC_RESPIRATION, GOBP_AEROBIC_RESPIRATION
- GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT, GOBP_MITOCHONDRIAL_TRANSMEMBRANE_TRANSPORT
- GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_NADH_TO_UBIQUINONE
- GOBP_ELECTRON_TRANSPORT_CHAIN, GOBP_ELECTRON_TRANSPORT_CHAIN
- GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GOBP_CRISTAE_FORMATION, GOBP_CRISTAE_FORMATION
- GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
- GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GOBP_POSITIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOBP_TRICARBOXYLIC_ACID_CYCLE, GOBP_TRICARBOXYLIC_ACID_CYCLE
- GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBOSE_PHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_DOPAMINE_METABOLIC_PROCESS, GOBP_DOPAMINE_METABOLIC_PROCESS
- GOBP_APOPTOTIC_MITOCHONDRIAL_CHANGES, GOBP_APOPTOTIC_MITOCHONDRIAL_CHANGES
- GOBP_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_TRANSPORTER_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_TRANSPORTER_ACTIVITY
- GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS, GOBP_TETRAPYRROLE_BIOSYNTHETIC_PROCESS
- GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT, GOBP_MITOCHONDRIAL_ATP_SYNTHESIS_COUPLED_PROTON_TRANSPORT
- GOBP_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_PORPHYRIN_CONTAINING_COMPOUND_METABOLIC_PROCESS
- GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS, GOBP_RIBONUCLEOSIDE_TRIPHOSPHATE_BIOSYNTHETIC_PROCESS
- GOBP_ATP_BIOSYNTHETIC_PROCESS, GOBP_ATP_BIOSYNTHETIC_PROCESS
- GOBP_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS, GOBP_NUCLEOSIDE_TRIPHOSPHATE_METABOLIC_PROCESS
- GOBP_PIGMENT_BIOSYNTHETIC_PROCESS, GOBP_PIGMENT_BIOSYNTHETIC_PROCESS
- GOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION, GOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION
- GOBP_HEME_METABOLIC_PROCESS, GOBP_HEME_METABOLIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_CATION_TRANSMEMBRANE_TRANSPORT
- GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_NEGATIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
- GOBP_DETOXIFICATION_OF_INORGANIC_COMPOUND, GOBP_DETOXIFICATION_OF_INORGANIC_COMPOUND
- GOBP_RESPONSE_TO_INCREASED_OXYGEN_LEVELS, GOBP_RESPONSE_TO_INCREASED_OXYGEN_LEVELS
- GOBP_CARDIOLIPIN_METABOLIC_PROCESS, GOBP_CARDIOLIPIN_METABOLIC_PROCESS
- GOBP_MITOCHONDRIAL_CALCIIUM_ION_TRANSMEMBRANE_TRANSPORT, GOBP_MITOCHONDRIAL_CALCIIUM_ION_TRANSMEMBRANE_TRANSPORT
- GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_CYTOCHROME_C_TO_OXYGEN, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_CYTOCHROME_C_TO_OXYGEN
- GOBP_DOPAMINE_TRANSPORT, GOBP_DOPAMINE_TRANSPORT
- GOBP_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS, GOBP_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_FATTY_ACID_BIOSYNTHETIC_PROCESS
- GOBP_UBIQUINONE_METABOLIC_PROCESS, GOBP_UBIQUINONE_METABOLIC_PROCESS
- GOBP_RESPONSE_TO_HYDROGEN_PEROXIDE, GOBP_RESPONSE_TO_HYDROGEN_PEROXIDE
- GOBP_PIGMENT_METABOLIC_PROCESS, GOBP_PIGMENT_METABOLIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GOBP_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
- GOBP_POSITIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS
- GOBP_PROTON_TRANSMEMBRANE_TRANSPORT, GOBP_PROTON_TRANSMEMBRANE_TRANSPORT
- GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_UBIQUINOL_TO_CYTOCHROME_C, GOBP_MITOCHONDRIAL_ELECTRON_TRANSPORT_UBIQUINOL_TO_CYTOCHROME_C
- GOBP_NUCLEOTIDE_TRANSPORT, GOBP_NUCLEOTIDE_TRANSPORT
- GOBP_POSITIVE_REGULATION_OF_CATION_CHANNEL_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_CATION_CHANNEL_ACTIVITY
- GOBP_NEURAL_NUCLEUS_DEVELOPMENT, GOBP_NEURAL_NUCLEUS_DEVELOPMENT