

MEMBRANE\_PROTEIN\_COMPLEX, GOCC\_INNER\_MITOCHONDRIAL\_MEMBRANE\_PROTEIN\_COMPLEX

- GOBP\_OXIDATIVE\_PHOSPHORYLATION, GOBP\_OXIDATIVE\_PHOSPHORYLATION
- GOBP\_CELLULAR\_RESPIRATION, GOBP\_CELLULAR\_RESPIRATION
- GOBP\_ATP\_SYNTHESIS\_COUPLED\_ELECTRON\_TRANSPORT, GOBP\_ATP\_SYNTHESIS\_COUPLED\_ELECTRON\_TRANSPORT
- GOBP\_PROTEIN\_TRANSMEMBRANE\_IMPORT\_INTO\_INTRACELLULAR\_ORGANELLE, GOBP\_PROTEIN\_TRANSMEMBRANE\_IMPORT\_INTO\_INTRACELLULAR\_ORGANELLE
- GOBP\_RESPIRATORY\_ELECTRON\_TRANSPORT\_CHAIN, GOBP\_RESPIRATORY\_ELECTRON\_TRANSPORT\_CHAIN
- GOCC\_RESPIRASOME, GOCC\_RESPIRASOME
- GOBP\_NADH\_DEHYDROGENASE\_COMPLEX\_ASSEMBLY, GOBP\_NADH\_DEHYDROGENASE\_COMPLEX\_ASSEMBLY
- GOCC\_OXIDOREDUCTASE\_COMPLEX, GOCC\_OXIDOREDUCTASE\_COMPLEX
- GOBP\_ELECTRON\_TRANSPORT\_CHAIN, GOBP\_ELECTRON\_TRANSPORT\_CHAIN
- GOBP\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRION, GOBP\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRION
- GOBP\_PROTEIN\_TARGETING\_TO\_MITOCHONDRION, GOBP\_PROTEIN\_TARGETING\_TO\_MITOCHONDRION
- GOCC\_INTRINSIC\_COMPONENT\_OF\_MITOCHONDRIAL\_MEMBRANE, GOCC\_INTRINSIC\_COMPONENT\_OF\_MITOCHONDRIAL\_MEMBRANE
- GOBP\_INNER\_MITOCHONDRIAL\_MEMBRANE\_ORGANIZATION, GOBP\_INNER\_MITOCHONDRIAL\_MEMBRANE\_ORGANIZATION
- GOBP\_MITOCHONDRIAL\_TRANSMEMBRANE\_TRANSPORT, GOBP\_MITOCHONDRIAL\_TRANSMEMBRANE\_TRANSPORT
- GOCC\_RESPIRATORY\_CHAIN\_COMPLEX, GOCC\_RESPIRATORY\_CHAIN\_COMPLEX
- GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_NAD\_P\_H, GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_NAD\_P\_H
- GOBP\_AEROBIC\_RESPIRATION, GOBP\_AEROBIC\_RESPIRATION
- GOCC\_ORGANELLE\_ENVELOPE\_LUMEN, GOCC\_ORGANELLE\_ENVELOPE\_LUMEN
- GOBP\_PROTEIN\_TRANSMEMBRANE\_TRANSPORT, GOBP\_PROTEIN\_TRANSMEMBRANE\_TRANSPORT
- GOBP\_MITOCHONDRIAL\_MEMBRANE\_ORGANIZATION, GOBP\_MITOCHONDRIAL\_MEMBRANE\_ORGANIZATION
- GOMF\_ELECTRON\_TRANSFER\_ACTIVITY, GOMF\_ELECTRON\_TRANSFER\_ACTIVITY
- GOBP\_MITOCHONDRIAL\_ELECTRON\_TRANSPORT\_NADH\_TO\_UBIQUINONE, GOBP\_MITOCHONDRIAL\_ELECTRON\_TRANSPORT\_NADH\_TO\_UBIQUINONE
- GOBP\_PROTEIN\_IMPORT\_INTO\_MITOCHONDRIAL\_MATRIX, GOBP\_PROTEIN\_IMPORT\_INTO\_MITOCHONDRIAL\_MATRIX
- GOMF\_NAD\_P\_H\_DEHYDROGENASE\_QUINONE\_ACTIVITY, GOMF\_NAD\_P\_H\_DEHYDROGENASE\_QUINONE\_ACTIVITY
- 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
- GOCC\_NADH\_DEHYDROGENASE\_COMPLEX, GOCC\_NADH\_DEHYDROGENASE\_COMPLEX
- GOCC\_INTRINSIC\_COMPONENT\_OF\_MITOCHONDRIAL\_INNER\_MEMBRANE, GOCC\_INTRINSIC\_COMPONENT\_OF\_MITOCHONDRIAL\_INNER\_MEMBRANE
- GOCC\_OUTER\_MITOCHONDRIAL\_MEMBRANE\_PROTEIN\_COMPLEX, GOCC\_OUTER\_MITOCHONDRIAL\_MEMBRANE\_PROTEIN\_COMPLEX
- GOBP\_RIBOSE\_PHOSPHATE\_BIOSYNTHETIC\_PROCESS, GOBP\_RIBOSE\_PHOSPHATE\_BIOSYNTHETIC\_PROCESS
- GOBP\_TRICARBOXYLIC\_ACID\_CYCLE, GOBP\_TRICARBOXYLIC\_ACID\_CYCLE
- GOCC\_TIM23\_MITOCHONDRIAL\_IMPORT\_INNER\_MEMBRANE\_TRANSLOCASE\_COMPLEX, GOCC\_TIM23\_MITOCHONDRIAL\_IMPORT\_INNER\_MEMBRANE\_TRANSLOCASE\_COMPLEX
- GOBP\_CRISTAE\_FORMATION, GOBP\_CRISTAE\_FORMATION
- GOMF\_PROTEIN\_TRANSPORTER\_ACTIVITY, GOMF\_PROTEIN\_TRANSPORTER\_ACTIVITY
- GOBP\_PROTEIN\_INSERTION\_INTO\_MEMBRANE, GOBP\_PROTEIN\_INSERTION\_INTO\_MEMBRANE
- GOMF\_MACROMOLECULE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_MACROMOLECULE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY
- GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRIAL\_MEMBRANE, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_MITOCHONDRIAL\_MEMBRANE
- GOMF\_PROTEIN\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_PROTEIN\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY
- GOMF\_AMIDE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_AMIDE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY
- GOCC\_SAM\_COMPLEX, GOCC\_SAM\_COMPLEX
- GOBP\_PROTEIN\_INSERTION\_INTO\_MITOCHONDRIAL\_INNER\_MEMBRANE, GOBP\_PROTEIN\_INSERTION\_INTO\_MITOCHONDRIAL\_INNER\_MEMBRANE
- GOMF\_PEPTIDE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_PEPTIDE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY
- GOMF\_MITOCHONDRION\_TARGETING\_SEQUENCE\_BINDING, GOMF\_MITOCHONDRION\_TARGETING\_SEQUENCE\_BINDING
- GOBP\_NUCLEOSIDE\_TRIPHOSPHATE\_BIOSYNTHETIC\_PROCESS, GOBP\_NUCLEOSIDE\_TRIPHOSPHATE\_BIOSYNTHETIC\_PROCESS
- GOMF\_2\_IRON\_2\_SULFUR\_CLUSTER\_BINDING, GOMF\_2\_IRON\_2\_SULFUR\_CLUSTER\_BINDING
- GOBP\_ATP\_SYNTHESIS\_COUPLED\_PROTON\_TRANSPORT, GOBP\_ATP\_SYNTHESIS\_COUPLED\_PROTON\_TRANSPORT
- GOBP\_RIBONUCLEOSIDE\_TRIPHOSPHATE\_BIOSYNTHETIC\_PROCESS, GOBP\_RIBONUCLEOSIDE\_TRIPHOSPHATE\_BIOSYNTHETIC\_PROCESS
- GOBP\_REGULATION\_OF\_ATPASE\_ACTIVITY, GOBP\_REGULATION\_OF\_ATPASE\_ACTIVITY
- GOCC\_CYTOCHROME\_COMPLEX, GOCC\_CYTOCHROME\_COMPLEX
- GOCC\_MICOS\_COMPLEX, GOCC\_MICOS\_COMPLEX
- GOCC\_PROTON\_TRANSPORTING\_TWO\_SECTOR\_ATPASE\_COMPLEX\_PROTON\_TRANSPORTING\_DOMAIN, GOCC\_PROTON\_TRANSPORTING\_TWO\_SECTOR\_ATPASE\_COMPLEX\_PROTON\_TRANSPORTING\_DOMAIN
- GOMF\_UNFOLDED\_PROTEIN\_BINDING, GOMF\_UNFOLDED\_PROTEIN\_BINDING
- GOBP\_MITOCHONDRIAL\_ATP\_SYNTHESIS\_COUPLED\_PROTON\_TRANSPORT, GOBP\_MITOCHONDRIAL\_ATP\_SYNTHESIS\_COUPLED\_PROTON\_TRANSPORT