

**GOCC\_MICROBODY, GOCC\_MICROBODY**

- GOBP\_PEROXISOME\_ORGANIZATION, GOBP\_PEROXISOME\_ORGANIZATION
- GOBP\_PEROXISOMAL\_TRANSPORT, GOBP\_PEROXISOMAL\_TRANSPORT
- GOMF\_FLAVIN\_ADENINE\_DINUCLEOTIDE\_BINDING, GOMF\_FLAVIN\_ADENINE\_DINUCLEOTIDE\_BINDING
- GOCC\_MICROBODY\_MEMBRANE, GOCC\_MICROBODY\_MEMBRANE
- GOBP\_MONOCARBOXYLIC\_ACID\_CATABOLIC\_PROCESS, GOBP\_MONOCARBOXYLIC\_ACID\_CATABOLIC\_PROCESS
- GOMF\_ISOMERASE\_ACTIVITY, GOMF\_ISOMERASE\_ACTIVITY
- GOBP\_ALPHA\_AMINO\_ACID\_METABOLIC\_PROCESS, GOBP\_ALPHA\_AMINO\_ACID\_METABOLIC\_PROCESS
- GOMF\_CARBOXYLIC\_ACID\_BINDING, GOMF\_CARBOXYLIC\_ACID\_BINDING
- GOBP\_THIOESTER\_BIOSYNTHETIC\_PROCESS, GOBP\_THIOESTER\_BIOSYNTHETIC\_PROCESS
- GOBP\_NUCLEOSIDE\_BISPHOSPHATE\_BIOSYNTHETIC\_PROCESS, GOBP\_NUCLEOSIDE\_BISPHOSPHATE\_BIOSYNTHETIC\_PROCESS
- GOBP\_FATTY\_ACID\_CATABOLIC\_PROCESS, GOBP\_FATTY\_ACID\_CATABOLIC\_PROCESS
- GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_CH\_OH\_GROUP\_OF\_DONORS, GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_CH\_OH\_GROUP\_OF\_DONORS
- GOBP\_ACETYL\_COA\_METABOLIC\_PROCESS, GOBP\_ACETYL\_COA\_METABOLIC\_PROCESS
- GOBP\_FATTY\_ACID\_BETA\_OXIDATION, GOBP\_FATTY\_ACID\_BETA\_OXIDATION
- GOBP\_LONG\_CHAIN\_FATTY\_ACID\_METABOLIC\_PROCESS, GOBP\_LONG\_CHAIN\_FATTY\_ACID\_METABOLIC\_PROCESS
- GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_ALKYL\_OR\_ARYL\_OTHER\_THAN\_METHYL\_GROUPS, GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_ALKYL\_OR\_ARYL\_OTHER\_THAN\_METHYL\_GROUPS
- GOBP\_NUCLEOSIDE\_BISPHOSPHATE\_METABOLIC\_PROCESS, GOBP\_NUCLEOSIDE\_BISPHOSPHATE\_METABOLIC\_PROCESS
- GOMF\_PRIMARY\_ACTIVE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_PRIMARY\_ACTIVE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY
- GOMF\_FAD\_BINDING, GOMF\_FAD\_BINDING
- GOBP\_AMINO\_ACID\_BETAINE\_METABOLIC\_PROCESS, GOBP\_AMINO\_ACID\_BETAINE\_METABOLIC\_PROCESS