

ME\_ORGANIZATION, GOBP\_PEROXISOME\_ORGANIZATION

- GOBP\_PEROXISOMAL\_TRANSPORT, GOBP\_PEROXISOMAL\_TRANSPORT
- GOCC\_MICROBODY\_MEMBRANE, GOCC\_MICROBODY\_MEMBRANE
- HP\_ORGANIC\_ACIDURIA, HP\_ORGANIC\_ACIDURIA
- HP\_AMINOACIDURIA, HP\_AMINOACIDURIA
- GOBP\_FATTY\_ACID\_CATABOLIC\_PROCESS, GOBP\_FATTY\_ACID\_CATABOLIC\_PROCESS
- GOBP\_THIOESTER\_METABOLIC\_PROCESS, GOBP\_THIOESTER\_METABOLIC\_PROCESS
- GOBP\_PEROXISOMAL\_MEMBRANE\_TRANSPORT, GOBP\_PEROXISOMAL\_MEMBRANE\_TRANSPORT
- GOBP\_VERY\_LONG\_CHAIN\_FATTY\_ACID\_METABOLIC\_PROCESS, GOBP\_VERY\_LONG\_CHAIN\_FATTY\_ACID\_METABOLIC\_PROCESS
- GOBP\_FATTY\_ACID\_BETA\_OXIDATION, GOBP\_FATTY\_ACID\_BETA\_OXIDATION
- GOBP\_NUCLEOSIDE\_BISPHOSPHATE\_METABOLIC\_PROCESS, GOBP\_NUCLEOSIDE\_BISPHOSPHATE\_METABOLIC\_PROCESS
- GOMF\_LYASE\_ACTIVITY, GOMF\_LYASE\_ACTIVITY
- GOBP\_ALPHA\_AMINO\_ACID\_CATABOLIC\_PROCESS, GOBP\_ALPHA\_AMINO\_ACID\_CATABOLIC\_PROCESS
- GOBP\_FATTY\_ACYL\_COA\_METABOLIC\_PROCESS, GOBP\_FATTY\_ACYL\_COA\_METABOLIC\_PROCESS
- HP\_ABNORMAL\_CIRCULATING\_DICARBOXYLIC\_ACID\_CONCENTRATION, HP\_ABNORMAL\_CIRCULATING\_DICARBOXYLIC\_ACID\_CONCENTRATION
- GOBP\_LONG\_CHAIN\_FATTY\_ACID\_METABOLIC\_PROCESS, GOBP\_LONG\_CHAIN\_FATTY\_ACID\_METABOLIC\_PROCESS
- GOBP\_PROTEIN\_IMPORT\_INTO\_PEROXISOME\_MATRIX, GOBP\_PROTEIN\_IMPORT\_INTO\_PEROXISOME\_MATRIX
- GOBP\_LIPID\_OXIDATION, GOBP\_LIPID\_OXIDATION
- HP\_ABNORMAL\_TISSUE\_METABOLITE\_CONCENTRATION, HP\_ABNORMAL\_TISSUE\_METABOLITE\_CONCENTRATION
- GOBP\_DICARBOXYLIC\_ACID\_CATABOLIC\_PROCESS, GOBP\_DICARBOXYLIC\_ACID\_CATABOLIC\_PROCESS
- HP\_VERY\_LONG\_CHAIN\_FATTY\_ACID\_ACCUMULATION, HP\_VERY\_LONG\_CHAIN\_FATTY\_ACID\_ACCUMULATION
- GOBP\_MONOCARBOXYLIC\_ACID\_CATABOLIC\_PROCESS, GOBP\_MONOCARBOXYLIC\_ACID\_CATABOLIC\_PROCESS
- GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_THE\_CH\_NH2\_GROUP\_OF\_DONORS, GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_THE\_CH\_NH2\_GROUP\_OF\_DONORS
- HP\_ABNORMAL\_CIRCULATING\_LONG\_CHAIN\_FATTY\_ACID\_CONCENTRATION, HP\_ABNORMAL\_CIRCULATING\_LONG\_CHAIN\_FATTY\_ACID\_CONCENTRATION
- GOBP\_CELLULAR\_AMINO\_ACID\_CATABOLIC\_PROCESS, GOBP\_CELLULAR\_AMINO\_ACID\_CATABOLIC\_PROCESS
- HP\_ABNORMAL\_CIRCULATING\_FATTY\_ACID\_CONCENTRATION, HP\_ABNORMAL\_CIRCULATING\_FATTY\_ACID\_CONCENTRATION
- HP\_BRUSHFIELD\_SPOTS, HP\_BRUSHFIELD\_SPOTS
- GOBP\_NUCLEOBASE\_CONTAINING\_SMALL\_MOLECULE\_BIOSYNTHETIC\_PROCESS, GOBP\_NUCLEOBASE\_CONTAINING\_SMALL\_MOLECULE\_BIOSYNTHETIC\_PROCESS
- GOBP\_FATTY\_ACID\_DERIVATIVE\_METABOLIC\_PROCESS, GOBP\_FATTY\_ACID\_DERIVATIVE\_METABOLIC\_PROCESS
- GOBP\_FATTY\_ACID\_DERIVATIVE\_CATABOLIC\_PROCESS, GOBP\_FATTY\_ACID\_DERIVATIVE\_CATABOLIC\_PROCESS
- HP\_ABNORMAL\_EPIPHYSEAL\_OSSIFICATION, HP\_ABNORMAL\_EPIPHYSEAL\_OSSIFICATION
- HP\_ABNORMALITY\_OF\_VITAMIN\_METABOLISM, HP\_ABNORMALITY\_OF\_VITAMIN\_METABOLISM
- HP\_DICARBOXYLIC\_ACIDURIA, HP\_DICARBOXYLIC\_ACIDURIA
- GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_CH\_OH\_GROUP\_OF\_DONORS, GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_CH\_OH\_GROUP\_OF\_DONORS
- GOBP\_MYD88\_INDEPENDENT\_TOLL\_LIKE\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_MYD88\_INDEPENDENT\_TOLL\_LIKE\_RECEPTOR\_SIGNALING\_PATHWAY
- GOBP\_OLEFINIC\_COMPOUND\_METABOLIC\_PROCESS, GOBP\_OLEFINIC\_COMPOUND\_METABOLIC\_PROCESS
- GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_THE\_CH\_CH\_GROUP\_OF\_DONORS\_NAD\_OR\_NADP\_AS\_ACCEPTOR, GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_THE\_CH\_CH\_GROUP\_OF\_DONORS\_NAD\_OR\_NADP\_AS\_ACCEPTOR
- HP\_HOMOCYSTINURIA, HP\_HOMOCYSTINURIA