

GENASE_ACTIVITY, GOMF_MONOOXYGENASE_ACTIVITY

- GOMF_STEROID_HYDROXYLASE_ACTIVITY, GOMF_STEROID_HYDROXYLASE_ACTIVITY
- GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS
- GOBP_REGULATION_OF_GUANYLATE_CYCLASE_ACTIVITY, GOBP_REGULATION_OF_GUANYLATE_CYCLASE_ACTIVITY
- GOBP_ALPHA_AMINO_ACID_METABOLIC_PROCESS, GOBP_ALPHA_AMINO_ACID_METABOLIC_PROCESS
- GOBP_CELLULAR_HORMONE_METABOLIC_PROCESS, GOBP_CELLULAR_HORMONE_METABOLIC_PROCESS
- GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_REDUCED_FLAVIN_OR_FLAVOPROTEIN_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_OF_OXYGEN, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_REDUCED_FLAVIN_OR_FLAVOPROTEIN_AS_ONE_DONOR
- GOBP_RETINOIC_ACID_METABOLIC_PROCESS, GOBP_RETINOIC_ACID_METABOLIC_PROCESS
- GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS, GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS
- GOBP_POSITIVE_REGULATION_OF_GUANYLATE_CYCLASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_GUANYLATE_CYCLASE_ACTIVITY
- GOMF_AROMATASE_ACTIVITY, GOMF_AROMATASE_ACTIVITY
- GOBP_EPOXYGENASE_P450_PATHWAY, GOBP_EPOXYGENASE_P450_PATHWAY
- GOBP_DRUG_METABOLIC_PROCESS, GOBP_DRUG_METABOLIC_PROCESS
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_OF_OXYGEN, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_PAIRED_DONORS_WITH_INCORPORATION_OR_REDUCTION_OF_MOLECULAR_OXYGEN_NAD_P_H_AS_ONE_DONOR_AND_INCORPORATION_OF_ONE_ATOM_OF_OXYGEN
- GOMF_RETINOIC_ACID_4_HYDROXYLASE_ACTIVITY, GOMF_RETINOIC_ACID_4_HYDROXYLASE_ACTIVITY
- GOBP_GLUCOCORTICOID_BIOSYNTHETIC_PROCESS, GOBP_GLUCOCORTICOID_BIOSYNTHETIC_PROCESS
- GOBP_LIPID_DIGESTION, GOBP_LIPID_DIGESTION
- GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS, GOBP_OLEFINIC_COMPOUND_METABOLIC_PROCESS
- GOMF_VITAMIN_BINDING, GOMF_VITAMIN_BINDING
- GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS, GOBP_INDOLALKYLAMINE_METABOLIC_PROCESS
- GOBP_AMINE_CATABOLIC_PROCESS, GOBP_AMINE_CATABOLIC_PROCESS
- GOMF_ARACHIDONIC_ACID_MONOOXYGENASE_ACTIVITY, GOMF_ARACHIDONIC_ACID_MONOOXYGENASE_ACTIVITY
- GOBP_ARACHIDONIC_ACID_METABOLIC_PROCESS, GOBP_ARACHIDONIC_ACID_METABOLIC_PROCESS
- GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GOBP_INDOLE_CONTAINING_COMPOUND_METABOLIC_PROCESS
- GOBP_REGULATION_OF_INTESTINAL_CHOLESTEROL_ABSORPTION, GOBP_REGULATION_OF_INTESTINAL_CHOLESTEROL_ABSORPTION
- GOBP_TYROSINE_METABOLIC_PROCESS, GOBP_TYROSINE_METABOLIC_PROCESS
- GOBP_DRUG_CATABOLIC_PROCESS, GOBP_DRUG_CATABOLIC_PROCESS
- GOBP_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS, GOBP_LONG_CHAIN_FATTY_ACID_BIOSYNTHETIC_PROCESS
- GOBP_STEROID_CATABOLIC_PROCESS, GOBP_STEROID_CATABOLIC_PROCESS