

D_BINDING, GOMF_NAD_BINDING

- GOCC_MICROBODY, GOCC_MICROBODY
- GOMF_LYASE_ACTIVITY, GOMF_LYASE_ACTIVITY
- GOBP_TRICARBOXYLIC_ACID_METABOLIC_PROCESS, GOBP_TRICARBOXYLIC_ACID_METABOLIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_PROTEIN_ACETYLATION, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_ACETYLATION
- GOBP_ALPHA_AMINO_ACID_METABOLIC_PROCESS, GOBP_ALPHA_AMINO_ACID_METABOLIC_PROCESS
- GOBP_SMAD_PROTEIN_COMPLEX_ASSEMBLY, GOBP_SMAD_PROTEIN_COMPLEX_ASSEMBLY
- GOBP_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS, GOBP_NUCLEOTIDE_SUGAR_BIOSYNTHETIC_PROCESS
- GOBP_NUCLEOTIDE_SUGAR_METABOLIC_PROCESS, GOBP_NUCLEOTIDE_SUGAR_METABOLIC_PROCESS
- GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS, GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS
- GOBP_ETHANOL_CATABOLIC_PROCESS, GOBP_ETHANOL_CATABOLIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION, GOBP_NEGATIVE_REGULATION_OF_PEPTIDYL_LYSINE_ACETYLATION
- GOMF_NADH_PYROPHOSPHATASE_ACTIVITY, GOMF_NADH_PYROPHOSPHATASE_ACTIVITY
- GOCC_MICROBODY_LUMEN, GOCC_MICROBODY_LUMEN
- GOBP_PRIMARY_ALCOHOL_METABOLIC_PROCESS, GOBP_PRIMARY_ALCOHOL_METABOLIC_PROCESS
- GOBP_CELLULAR_AMINO_ACID_CATABOLIC_PROCESS, GOBP_CELLULAR_AMINO_ACID_CATABOLIC_PROCESS
- GOBP_TERPENOID_METABOLIC_PROCESS, GOBP_TERPENOID_METABOLIC_PROCESS
- GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS_IN_LINEAR_AMIDES, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS_IN_LINEAR_AMIDES
- GOBP_DICARBOXYLIC_ACID_METABOLIC_PROCESS, GOBP_DICARBOXYLIC_ACID_METABOLIC_PROCESS
- GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_CARBON_NITROGEN_BUT_NOT_PEPTIDE_BONDS
- GOBP_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHOSPHATE, GOBP_GLYCOLYTIC_PROCESS_THROUGH_FRUCTOSE_6_PHOSPHATE
- GOBP_VALINE_METABOLIC_PROCESS, GOBP_VALINE_METABOLIC_PROCESS
- GOMF_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS, GOMF_HYDROLASE_ACTIVITY_ACTING_ON_ETHER_BONDS
- GOBP_NAD_BIOSYNTHESIS_VIA_NICOTINAMIDE_RIBOSIDE_SALVAGE_PATHWAY, GOBP_NAD_BIOSYNTHESIS_VIA_NICOTINAMIDE_RIBOSIDE_SALVAGE_PATHWAY
- GOBP_NADH_OXIDATION, GOBP_NADH_OXIDATION
- GOCC_TRICARBOXYLIC_ACID_CYCLE_ENZYME_COMPLEX, GOCC_TRICARBOXYLIC_ACID_CYCLE_ENZYME_COMPLEX
- GOBP_PURINE_NUCLEOSIDE_CATABOLIC_PROCESS, GOBP_PURINE_NUCLEOSIDE_CATABOLIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_HISTONE_ACETYLATION, GOBP_NEGATIVE_REGULATION_OF_HISTONE_ACETYLATION
- GOCC_PYRUVATE_DEHYDROGENASE_COMPLEX, GOCC_PYRUVATE_DEHYDROGENASE_COMPLEX
- GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_OTHER_NITROGENOUS_COMPOUNDS_AS_DONORS, GOMF_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_OTHER_NITROGENOUS_COMPOUNDS_AS_DONORS
- GOBP_ESTROGEN_BIOSYNTHETIC_PROCESS, GOBP_ESTROGEN_BIOSYNTHETIC_PROCESS
- GOMF_NADPLUS_BINDING, GOMF_NADPLUS_BINDING
- GOBP_PRIMARY_ALCOHOL_CATABOLIC_PROCESS, GOBP_PRIMARY_ALCOHOL_CATABOLIC_PROCESS