

TE_METABOLIC_PROCESS, GO_TETRAHYDROFOLATE_METABOLIC_PROCESS

- GO_L_SERINE_METABOLIC_PROCESS, GO_L_SERINE_METABOLIC_PROCESS
- GO_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GO_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GO_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS, GO_PURINE_NUCLEOBASE_BIOSYNTHETIC_PROCESS
- GO_VITAMIN_B6_BINDING, GO_VITAMIN_B6_BINDING
- GO_PURINE_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS, GO_PURINE_NUCLEOSIDE_MONOPHOSPHATE_BIOSYNTHETIC_PROCESS
- GO_FOLIC_ACID_METABOLIC_PROCESS, GO_FOLIC_ACID_METABOLIC_PROCESS
- GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS
- GO_FOLIC_ACID_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_FOLIC_ACID_CONTAINING_COMPOUND_METABOLIC_PROCESS
- GO_PYRIMIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_PYRIMIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
- GO_CELLULAR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_CELLULAR_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GO_NADPH_BINDING, GO_NADPH_BINDING
- GO_CARBON_CARBON_LYASE_ACTIVITY, GO_CARBON_CARBON_LYASE_ACTIVITY
- GO_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GO_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GO_PYRIMIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GO_PYRIMIDINE_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
- GO_CELLULAR_MODIFIED_AMINO_ACID_CATABOLIC_PROCESS, GO_CELLULAR_MODIFIED_AMINO_ACID_CATABOLIC_PROCESS
- GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS
- GO_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GO_SULFUR_AMINO_ACID_METABOLIC_PROCESS
- GO_TETRAHYDROFOLATE_INTERCONVERSION, GO_TETRAHYDROFOLATE_INTERCONVERSION
- GO_PURINE_NUCLEOBASE_METABOLIC_PROCESS, GO_PURINE_NUCLEOBASE_METABOLIC_PROCESS
- GO_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS, GO_DEOXYRIBONUCLEOSIDE_MONOPHOSPHATE_METABOLIC_PROCESS
- GO_IMP_BIOSYNTHETIC_PROCESS, GO_IMP_BIOSYNTHETIC_PROCESS