

CID_BIOSYNTHETIC_PROCESS, GOBP_ALPHA_AMINO_ACID_BIOSYNTHETIC_PROCESS

- GOMF_LYASE_ACTIVITY, GOMF_LYASE_ACTIVITY
- GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS
- HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_PROTEINOGENIC_AMINO_ACID_CONCENTRATION
- GOBP_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_SERINE_FAMILY_AMINO_ACID_BIOSYNTHETIC_PROCESS
- HP_ABNORMAL_CIRCULATING_SERINE_FAMILY_AMINO_ACID_CONCENTRATION, HP_ABNORMAL_CIRCULATING_SERINE_FAMILY_AMINO_ACID_CONCENTRATION
- GOBP_ANION_HOMEOSTASIS, GOBP_ANION_HOMEOSTASIS
- GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS, GOBP_BRANCHED_CHAIN_AMINO_ACID_METABOLIC_PROCESS
- GOBP_2_OXOGLUTARATE_METABOLIC_PROCESS, GOBP_2_OXOGLUTARATE_METABOLIC_PROCESS
- GOMF_HYDRO_LYASE_ACTIVITY, GOMF_HYDRO_LYASE_ACTIVITY
- GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS, GOBP_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
- GOMF_FLAVIN_ADENINE_DINUCLEOTIDE_BINDING, GOMF_FLAVIN_ADENINE_DINUCLEOTIDE_BINDING
- GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS, GOBP_SULFUR_AMINO_ACID_METABOLIC_PROCESS
- GOBP_PHENOL_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS, GOBP_PHENOL_CONTAINING_COMPOUND_BIOSYNTHETIC_PROCESS
- GOBP_L_SERINE_METABOLIC_PROCESS, GOBP_L_SERINE_METABOLIC_PROCESS
- GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS, GOBP_SULFUR_AMINO_ACID_BIOSYNTHETIC_PROCESS
- GOBP_PROTEIN_HYDROXYLATION, GOBP_PROTEIN_HYDROXYLATION