

GO_PROTEIN_TARGETING_TO_MITOCHONDRION, GO_PROTEIN_TARGETING_TO_MITOCHONDRION

- GO_REGULATION_OF_MACROAUTOPHAGY, GO_REGULATION_OF_MACROAUTOPHAGY
- GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT, GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT
- GO_VACUOLE_ORGANIZATION, GO_VACUOLE_ORGANIZATION
- GO_UBIQUITIN_LIKE_PROTEIN_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_BINDING
- GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
- GO_VACUOLAR_TRANSPORT, GO_VACUOLAR_TRANSPORT
- GO_AUTOPHAGOSOME_ORGANIZATION, GO_AUTOPHAGOSOME_ORGANIZATION
- GO_UBIQUITIN_BINDING, GO_UBIQUITIN_BINDING
- GO_PROTEIN_K48_LINKED_UBIQUITINATION, GO_PROTEIN_K48_LINKED_UBIQUITINATION
- GO_PROTEIN_K11_LINKED_UBIQUITINATION, GO_PROTEIN_K11_LINKED_UBIQUITINATION
- GO_AUTOPHAGY_OF_MITOCHONDRION, GO_AUTOPHAGY_OF_MITOCHONDRION
- GO_ORGANELLE_FUSION, GO_ORGANELLE_FUSION
- GO_REGULATION_OF_VACUOLE_ORGANIZATION, GO_REGULATION_OF_VACUOLE_ORGANIZATION
- GO_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_ATPASE_ACTIVITY
- GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRIAL_MEMBRANE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRIAL_MEMBRANE
- GO_MITOPHAGY, GO_MITOPHAGY
- GO_SIGNAL_SEQUENCE_BINDING, GO_SIGNAL_SEQUENCE_BINDING
- GO_PEROXISOME_ORGANIZATION, GO_PEROXISOME_ORGANIZATION
- GO_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION, GO_POSITIVE_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION
- GO_ACTOMYOSIN, GO_ACTOMYOSIN
- GO_REGULATION_OF_PROTEIN_TARGETING, GO_REGULATION_OF_PROTEIN_TARGETING
- GO_OUTER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX, GO_OUTER_MITOCHONDRIAL_MEMBRANE_PROTEIN_COMPLEX
- GO_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION, GO_REGULATION_OF_PROTEIN_TARGETING_TO_MITOCHONDRION
- GO_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_INNER_MEMBRANE, GO_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_INNER_MEMBRANE
- GO_PROTEIN_AUTOUBIQUITINATION, GO_PROTEIN_AUTOUBIQUITINATION
- GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION, GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION
- GO_CD40_RECEPTOR_COMPLEX, GO_CD40_RECEPTOR_COMPLEX
- GO_PROTEIN_TRANSMEMBRANE_TRANSPORT, GO_PROTEIN_TRANSMEMBRANE_TRANSPORT
- GO_NEGATIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_ENDOPLASMIC_RETICULUM_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
- GO_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE, GO_COPII_COATED_ER_TO_GOLGI_TRANSPORT_VESICLE
- GO_CHAPERONE_MEDIATED_PROTEIN_TRANSPORT, GO_CHAPERONE_MEDIATED_PROTEIN_TRANSPORT
- GO_PROTEIN_MATURATION_BY_PROTEIN_FOLDING, GO_PROTEIN_MATURATION_BY_PROTEIN_FOLDING
- GO_SELECTIVE_AUTOPHAGY, GO_SELECTIVE_AUTOPHAGY
- GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY, GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY