

_LIKE_PROTEIN_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_BINDING

- GO_UBIQUITIN_BINDING, GO_UBIQUITIN_BINDING
- GO_SCF_UBIQUITIN_LIGASE_COMPLEX, GO_SCF_UBIQUITIN_LIGASE_COMPLEX
- GO_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GO_ERAD_PATHWAY, GO_ERAD_PATHWAY
- GO_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT, GO_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT
- GO_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY, GO_STEROID_HORMONE_MEDIATED_SIGNALING_PATHWAY
- GO_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GO_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
- GO_PROTEIN_K48_LINKED_UBIQUITINATION, GO_PROTEIN_K48_LINKED_UBIQUITINATION
- GO_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GO_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY
- GO_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
- GO_HISTONE_UBIQUITINATION, GO_HISTONE_UBIQUITINATION
- GO_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY, GO_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY
- GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION, GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION
- GO_CORTICOSTEROID_RECEPTOR_SIGNALING_PATHWAY, GO_CORTICOSTEROID_RECEPTOR_SIGNALING_PATHWAY
- GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
- GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_OXIDATIVE_STRESS
- GO_HISTONE_H2A_UBIQUITINATION, GO_HISTONE_H2A_UBIQUITINATION
- GO_PROTEIN_AUTOUBIQUITINATION, GO_PROTEIN_AUTOUBIQUITINATION
- GO_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GO_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
- GO_HSP70_PROTEIN_BINDING, GO_HSP70_PROTEIN_BINDING
- GO_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY, GO_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY
- GO_RESPONSE_TO_MITOCHONDRIAL_DEPOLARISATION, GO_RESPONSE_TO_MITOCHONDRIAL_DEPOLARISATION
- GO_PROTEIN_DEACETYLASE_ACTIVITY, GO_PROTEIN_DEACETYLASE_ACTIVITY
- GO_AUTOPHAGY_OF_MITOCHONDRION, GO_AUTOPHAGY_OF_MITOCHONDRION
- GO_POSITIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_SELECTIVE_AUTOPHAGY, GO_SELECTIVE_AUTOPHAGY
- GO_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, GO_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
- GO_RESPONSE_TO_MISFOLDED_PROTEIN, GO_RESPONSE_TO_MISFOLDED_PROTEIN
- GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION_IN_RESPONSE_TO_MITOCHONDRIAL_DEPOLARIZATION, GO_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION_IN_RESPONSE_TO_MITOCHONDRIAL_DEPOLARIZATION
- GO_REGULATION_OF_ERAD_PATHWAY, GO_REGULATION_OF_ERAD_PATHWAY
- GO_REGULATION_OF_HISTONE_UBIQUITINATION, GO_REGULATION_OF_HISTONE_UBIQUITINATION
- GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
- GO_REGULATION_OF_PROTEIN_TARGETING, GO_REGULATION_OF_PROTEIN_TARGETING
- GO_EXTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE, GO_EXTRINSIC_COMPONENT_OF_ENDOPLASMIC_RETICULUM_MEMBRANE
- GO_NEGATIVE_REGULATION_OF_ERAD_PATHWAY, GO_NEGATIVE_REGULATION_OF_ERAD_PATHWAY
- GO_SUMO_TRANSFERASE_ACTIVITY, GO_SUMO_TRANSFERASE_ACTIVITY
- GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING