

_LIKE_PROTEIN_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_BINDING

- GO_UBIQUITIN_BINDING, GO_UBIQUITIN_BINDING
- GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
- GO_ERAD_PATHWAY, GO_ERAD_PATHWAY
- GO_PEPTIDASE_COMPLEX, GO_PEPTIDASE_COMPLEX
- GO_ENDOPEPTIDASE_COMPLEX, GO_ENDOPEPTIDASE_COMPLEX
- 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_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GO_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
- GO_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GO_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
- GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
- GO_MULTIVESICULAR_BODY_SORTING_PATHWAY, GO_MULTIVESICULAR_BODY_SORTING_PATHWAY
- GO_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY, GO_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY
- GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY_IN_RESPONSE_TO_VIRUS, GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY_IN_RESPONSE_TO_VIRUS
- GO_ENDOSOME_ORGANIZATION, GO_ENDOSOME_ORGANIZATION
- GO_CULLIN_FAMILY_PROTEIN_BINDING, GO_CULLIN_FAMILY_PROTEIN_BINDING
- GO_PROTEIN_K63_LINKED_DEUBIQUITINATION, GO_PROTEIN_K63_LINKED_DEUBIQUITINATION
- GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
- GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_BINDING
- GO_HISTONE_UBIQUITINATION, GO_HISTONE_UBIQUITINATION
- GO_POSITIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL, GO_POSITIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL
- GO_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS, GO_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS
- GO_MULTI_ORGANISM_CELLULAR_PROCESS, GO_MULTI_ORGANISM_CELLULAR_PROCESS
- GO_PHOSPHOPROTEIN_BINDING, GO_PHOSPHOPROTEIN_BINDING
- GO_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GO_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
- GO_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY, GO_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY
- GO_HISTONE_H2A_UBIQUITINATION, GO_HISTONE_H2A_UBIQUITINATION
- GO_NEGATIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
- GO_PHAGOPHORE_ASSEMBLY_SITE, GO_PHAGOPHORE_ASSEMBLY_SITE
- GO_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY, GO_ANDROGEN_RECEPTOR_SIGNALING_PATHWAY
- GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
- GO_EXTRACELLULAR_VESICLE_BIOGENESIS, GO_EXTRACELLULAR_VESICLE_BIOGENESIS
- GO_PROTEIN_MONOUBIQUITINATION, GO_PROTEIN_MONOUBIQUITINATION
- GO_ACTIVATION_OF_MAPK_ACTIVITY, GO_ACTIVATION_OF_MAPK_ACTIVITY
- GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS
- GO_REGULATION_OF_VIRAL_INDUCED_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GO_REGULATION_OF_VIRAL_INDUCED_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
- GO_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY, GO_INTRACELLULAR_STEROID_HORMONE_RECEPTOR_SIGNALING_PATHWAY
- GO_PROTEIN_K48_LINKED_UBIQUITINATION, GO_PROTEIN_K48_LINKED_UBIQUITINATION
- GO_RESPONSE_TO_MISFOLDED_PROTEIN, GO_RESPONSE_TO_MISFOLDED_PROTEIN
- GO_NUCLEAR_ENVELOPE_REASSEMBLY, GO_NUCLEAR_ENVELOPE_REASSEMBLY
- GO_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT, GO_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT
- GO_RIG_I_SIGNALING_PATHWAY, GO_RIG_I_SIGNALING_PATHWAY
- GO_SCF_UBIQUITIN_LIGASE_COMPLEX, GO_SCF_UBIQUITIN_LIGASE_COMPLEX
- GO_CYSTEINE_TYPE_ENDOPEPTIDASE_REGULATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GO_CYSTEINE_TYPE_ENDOPEPTIDASE_REGULATOR_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS