

N_LIKE_PROTEIN_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING

- GOCC_ENDOPEPTIDASE_COMPLEX, GOCC_ENDOPEPTIDASE_COMPLEX
- GOMF_UBIQUITIN_BINDING, GOMF_UBIQUITIN_BINDING
- GOCC_PEPTIDASE_COMPLEX, GOCC_PEPTIDASE_COMPLEX
- GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX
- GOBP_NEGATIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS
- GOBP_AUTOPHAGY_OF_MITOCHONDRION, GOBP_AUTOPHAGY_OF_MITOCHONDRION
- GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
- GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
- GOMF_HEAT_SHOCK_PROTEIN_BINDING, GOMF_HEAT_SHOCK_PROTEIN_BINDING
- GOBP_VACUOLAR_TRANSPORT, GOBP_VACUOLAR_TRANSPORT
- GOBP_POSITIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING, GOBP_POSITIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING
- GOBP_PROTEIN_DEGLYCOSYLATION, GOBP_PROTEIN_DEGLYCOSYLATION
- GOCC_AUTOPHAGOSOME, GOCC_AUTOPHAGOSOME
- HP_RIGIDITY, HP_RIGIDITY
- GOBP_ENDOSOME_ORGANIZATION, GOBP_ENDOSOME_ORGANIZATION
- GOBP_POSITIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY
- GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GOBP_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY
- GOMF_PROTEIN_DEACETYLASE_ACTIVITY, GOMF_PROTEIN_DEACETYLASE_ACTIVITY
- GOMF_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY, GOMF_UBIQUITIN_LIKE_PROTEIN_SPECIFIC_PROTEASE_ACTIVITY
- HP_ABNORMAL_CIRCULATING_IGG_LEVEL, HP_ABNORMAL_CIRCULATING_IGG_LEVEL
- GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS, GOBP_NEGATIVE_REGULATION_OF_PROTEIN_CATABOLIC_PROCESS
- GOBP_LYSOSOMAL_TRANSPORT, GOBP_LYSOSOMAL_TRANSPORT
- GOBP_ACTIVATION_OF_MAPK_ACTIVITY, GOBP_ACTIVATION_OF_MAPK_ACTIVITY
- GOBP_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GOBP_HISTONE_DEACETYLATION, GOBP_HISTONE_DEACETYLATION
- GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT, GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT
- GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GOBP_POSITIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
- GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOBP_ACTIVATION_OF_CYSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
- GOMF_CHAPERONE_BINDING, GOMF_CHAPERONE_BINDING
- GOMF_SUMO_BINDING, GOMF_SUMO_BINDING
- GOMF_PHOSPHOPROTEIN_BINDING, GOMF_PHOSPHOPROTEIN_BINDING
- GOBP_NUCLEAR_ENVELOPE_REASSEMBLY, GOBP_NUCLEAR_ENVELOPE_REASSEMBLY
- GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS, GOBP_EXTRACELLULAR_VESICLE_BIOGENESIS
- GOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION, GOBP_REGULATION_OF_PROTEIN_DEUBIQUITINATION
- GOBP_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GOBP_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
- GOBP_REGULATION_OF_ERAD_PATHWAY, GOBP_REGULATION_OF_ERAD_PATHWAY
- GOBP_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GOBP_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
- GOBP_ENDOPLASMIC_RETICULUM_MANNOSE_TRIMMING, GOBP_ENDOPLASMIC_RETICULUM_MANNOSE_TRIMMING
- GOBP_SELECTIVE_AUTOPHAGY, GOBP_SELECTIVE_AUTOPHAGY
- GOMF_DEACETYLASE_ACTIVITY, GOMF_DEACETYLASE_ACTIVITY
- GOBP_JNK_CASCADE, GOBP_JNK_CASCADE
- GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION_IN_RESPONSE_TO_MITOCHONDRIAL_DEPOLARIZATION, GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION_IN_RESPONSE_TO_MITOCHONDRIAL_DEPOLARIZATION
- GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY