

TOLL LIKE PROTEIN BINDING, GOMF_UBIQUITIN LIKE PROTEIN BINDING

- GOBP_PROTEIN_K48_LINKED_UBIQUITINATION, GOBP_PROTEIN_K48_LINKED_UBIQUITINATION
- GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX
- GOMF_UBIQUITIN_BINDING, GOMF_UBIQUITIN_BINDING
- GOBP_VACUOLAR_TRANSPORT, GOBP_VACUOLAR_TRANSPORT
- GOBP_PROTEIN_MONOUBIQUITINATION, GOBP_PROTEIN_MONOUBIQUITINATION
- GOBP_ERAD_PATHWAY, GOBP_ERAD_PATHWAY
- GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION
- GOCC_AUTOPHAGOSOME, GOCC_AUTOPHAGOSOME
- GOBP_ENDOSOME_ORGANIZATION, GOBP_ENDOSOME_ORGANIZATION
- GOBP_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_PROTEIN_LOCALIZATION_TO_VACUOLE
- GOBP_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM, GOBP_PROTEIN_EXIT_FROM_ENDOPLASMIC_RETICULUM
- GOBP_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GOBP_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
- GOBP_PROTEIN_DEGLYCOSYLATION, GOBP_PROTEIN_DEGLYCOSYLATION
- GOMF_PROTEIN_SELF_ASSOCIATION, GOMF_PROTEIN_SELF_ASSOCIATION
- GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
- GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT, GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT
- GOBP_HISTONE_H2A_UBIQUITINATION, GOBP_HISTONE_H2A_UBIQUITINATION
- HP_FASCICULATIONS, HP_FASCICULATIONS
- GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
- GOBP_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOBP_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
- GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT
- GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE
- HP_ASTROCYTOSIS, HP_ASTROCYTOSIS
- GOMF_SMAD_BINDING, GOMF_SMAD_BINDING
- GOBP_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY, GOBP_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY
- GOBP_REGULATION_OF_ERAD_PATHWAY, GOBP_REGULATION_OF_ERAD_PATHWAY
- GOBP_MYD88_DEPENDENT_TOLL LIKE RECEPTOR SIGNALING_PATHWAY, GOBP_MYD88_DEPENDENT_TOLL LIKE RECEPTOR SIGNALING_PATHWAY
- GOBP_NUCLEAR_ENVELOPE_REASSEMBLY, GOBP_NUCLEAR_ENVELOPE_REASSEMBLY
- GOMF_CULLIN_FAMILY_PROTEIN_BINDING, GOMF_CULLIN_FAMILY_PROTEIN_BINDING
- GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
- GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY
- GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT
- HP_BONE_MARROW_HYPERCELLULARITY, HP_BONE_MARROW_HYPERCELLULARITY
- GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GOBP_NEGATIVE_REGULATION_OF_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
- GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION, GOBP_PROTEIN_K63_LINKED_DEUBIQUITINATION
- GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS, GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS
- GOCC_SCF_UBIQUITIN_LIGASE_COMPLEX, GOCC_SCF_UBIQUITIN_LIGASE_COMPLEX
- GOBP_REGULATION_OF_MITOTIC_CENTROSOME_SEPARATION, GOBP_REGULATION_OF_MITOTIC_CENTROSOME_SEPARATION
- GOMF_DEAD_H_BOX_RNA_HELICASE_BINDING, GOMF_DEAD_H_BOX_RNA_HELICASE_BINDING
- GOBP_PROTEIN_DEMANNOSYLATION, GOBP_PROTEIN_DEMANNOSYLATION
- GOBP_ENDOPLASMIC_RETICULUM_MANNOSE_TRIMMING, GOBP_ENDOPLASMIC_RETICULUM_MANNOSE_TRIMMING
- GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
- GOBP_LIPID_DROPLET_ORGANIZATION, GOBP_LIPID_DROPLET_ORGANIZATION
- HP_ECHOLALIA, HP_ECHOLALIA
- GOMF_CYSSTEINE_TYPE_PEPTIDASE_ACTIVITY, GOMF_CYSSTEINE_TYPE_PEPTIDASE_ACTIVITY