GOBP_VACUOLAR_TRANSPORT, GOBP_VACUOLAR_TRANSPORT
GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PROTEIN_CONTAINING_COMPLEX
GOBP_PROTEIN_K48_LINKED_UBIQUITINATION, GOBP_PROTEIN_K48_LINKED_UBIQUITINATION
GOMF_UBIQUITIN_BINDING, GOMF_UBIQUITIN_BINDING
GOCC_ENDOPEPTIDASE_COMPLEX, GOCC_ENDOPEPTIDASE_COMPLEX
GOCC_PEPTIDASE_COMPLEX, GOCC_PEPTIDASE_COMPLEX
GOBP_PROTEIN_MONOUBIQUITINATION, GOBP_PROTEIN_MONOUBIQUITINATION
GOBP_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
GOCC_AUTOPHAGOSOME, GOCC_AUTOPHAGOSOME
GOBP_ERAD_PATHWAY, GOBP_ERAD_PATHWAY
GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION
GOBP_ENDOSOME_ORGANIZATION, GOBP_ENDOSOME_ORGANIZATION
GOBP_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GOBP_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
GOBP_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_PROTEIN_LOCALIZATION_TO_VACUOLE
GOBP_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GOBP_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
GOBP_PROTEIN_DEGLYCOSYLATION, GOBP_PROTEIN_DEGLYCOSYLATION
GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE, GOBP_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_VACUOLE
GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT, GOBP_ENDOPLASMIC_RETICULUM_TO_CYTOSOL_TRANSPORT
GOBP_HISTONE_H2A_UBIQUITINATION, GOBP_HISTONE_H2A_UBIQUITINATION
GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_ENDOSOME_TRANSPORT_VIA_MULTIVESICULAR_BODY_SORTING_PATHWAY
GOMF_SMAD_BINDING, GOMF_SMAD_BINDING
GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT, GOCC_ENDOPLASMIC_RETICULUM_QUALITY_CONTROL_COMPARTMENT
GOBP_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY, GOBP_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEP
GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT, GOBP_LATE_ENDOSOME_TO_VACUOLE_TRANSPORT
GOBP_REGULATION_OF_ERAD_PATHWAY, GOBP_REGULATION_OF_ERAD_PATHWAY
GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY_IN_RESPONSE_TO_VIRUS, GOBP_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY_IN_RESPO
GOBP_PROTEIN_K63_LINKED_UBIQUITINATION, GOBP_PROTEIN_K63_LINKED_UBIQUITINATION
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
GOMF_PROTEIN_SELF_ASSOCIATION, GOMF_PROTEIN_SELF_ASSOCIATION
GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY, GOBP_MULTIVESICULAR_BODY_SORTING_PATHWAY
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
GOMF_OMEGA_PEPTIDASE_ACTIVITY, GOMF_OMEGA_PEPTIDASE_ACTIVITY
GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS, GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEIN
GOBP_NEGATIVE_REGULATION_OF_CELL_GROWTH, GOBP_NEGATIVE_REGULATION_OF_CELL_GROWTH
GOBP_RECEPTOR_CATABOLIC_PROCESS, GOBP_RECEPTOR_CATABOLIC_PROCESS
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_PA
GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_SIGNALING_RECEPTOR_ACTIVITY
GOBP_LIPID_DROPLET_ORGANIZATION, GOBP_LIPID_DROPLET_ORGANIZATION
GOBP_REGULATION_OF_VASCULAR_WOUND_HEALING, GOBP_REGULATION_OF_VASCULAR_WOUND_HEALING
GOMF_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY, GOMF_CYSTEINE_TYPE_PEPTIDASE_ACTIVITY
GOCC_SMAD_PROTEIN_COMPLEX, GOCC_SMAD_PROTEIN_COMPLEX
GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GOMF_K63_LINKED_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING

N_LIKE_PROTEIN_BINDING, GOMF_UBIQUITIN_LIKE_PROTEIN_BINDING