

UBIQUITIN_LIGASE_COMPLEX, GOCC_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX

- GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GOBP_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
- GOBP_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GOCC_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX
- GOBP_ERAD_PATHWAY, GOBP_ERAD_PATHWAY
- GOBP_POSITIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL, GOBP_POSITIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CO
- GOBP_PROTEIN_MONOUBIQUITINATION, GOBP_PROTEIN_MONOUBIQUITINATION
- GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION
- GOBP_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GOBP_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY, GOBP_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
- GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY, GOBP_UBIQUITIN_DEPENDENT_ERAD_PATHWAY
- GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL3_RING_UBIQUITIN_LIGASE_COMPLEX
- GOBP_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CAT
- GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING, GOBP_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING
- GOBP_POSITIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CAT
- GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GOBP_PROTEIN_K48_LINKED_UBIQUITINATION, GOBP_PROTEIN_K48_LINKED_UBIQUITINATION
- GOBP_POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_CELLULAR_PROTEIN_CATABOLIC_PROCESS
- GOBP_PHOTOPERIODISM, GOBP_PHOTOPERIODISM
- GOCC_SCF_UBIQUITIN_LIGASE_COMPLEX, GOCC_SCF_UBIQUITIN_LIGASE_COMPLEX
- GOBP_PROTEIN_AUTOUBIQUITINATION, GOBP_PROTEIN_AUTOUBIQUITINATION
- GOMF_PROTEIN_PHOSPHATASE_BINDING, GOMF_PROTEIN_PHOSPHATASE_BINDING
- GOCC_CUL4A_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GOCC_CUL4A_RING_E3_UBIQUITIN_LIGASE_COMPLEX
- GOBP_REGULATION_OF_TOR_SIGNALING, GOBP_REGULATION_OF_TOR_SIGNALING
- GOBP_PROTEIN_DESTABILIZATION, GOBP_PROTEIN_DESTABILIZATION
- GOCC_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX, GOCC_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX
- GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTEINS, GOBP_PROTEIN_QUALITY_CONTROL_FOR_MISFOLDED_OR_INCOMPLETELY_SYNTHESIZED_PROTI
- GOBP_NEGATIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION, GOBP_NEGATIVE_REGULATION_OF_SMALL_GTPASE_MEDIATED_SIGNAL_TRANSDUCTION
- GOBP_RESPONSE_TO_MISFOLDED_PROTEIN, GOBP_RESPONSE_TO_MISFOLDED_PROTEIN
- GOMF_CULLIN_FAMILY_PROTEIN_BINDING, GOMF_CULLIN_FAMILY_PROTEIN_BINDING
- GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION
- GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK, GOBP_ENTRAINMENT_OF_CIRCADIAN_CLOCK
- GOBP_POSITIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION, GOBP_POSITIVE_REGULATION_OF_CELL_MORPHOGENESIS_INVOLVED_IN_DIFFERENTIATION
- GOBP_HISTONE_UBIQUITINATION, GOBP_HISTONE_UBIQUITINATION
- GOBP_PROTEIN_K63_LINKED_UBIQUITINATION, GOBP_PROTEIN_K63_LINKED_UBIQUITINATION
- GOBP_PROTEIN_HOMOOIGOMERIZATION, GOBP_PROTEIN_HOMOOIGOMERIZATION
- GOBP_NEURON_DEATH_IN_RESPONSE_TO_OXIDATIVE_STRESS, GOBP_NEURON_DEATH_IN_RESPONSE_TO_OXIDATIVE_STRESS
- GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_PROTEIN_TRANSFERASE_ACTIVITY, GOBP_POSITIVE_REGULATION_OF_UBIQUITIN_PROTEIN_TRANSFERASE_ACTIVITY
- GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION, GOBP_REGULATION_OF_AUTOPHAGY_OF_MITOCHONDRION