

UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_PROTEIN_UBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS

- GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
- GO_UBIQUITIN_LIKE_PROTEIN_LIGASE_ACTIVITY, GO_UBIQUITIN_LIKE_PROTEIN_LIGASE_ACTIVITY
- GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_UBIQUITIN_LIKE_PROTEIN_BINDING, GO_UBIQUITIN_LIKE_PROTEIN_BINDING
- GO_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
- KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS, KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS
- GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT
- GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
- KEGG_ENDOCYTOSIS, KEGG_ENDOCYTOSIS
- GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
- GO_RESPONSE_TO_UV, GO_RESPONSE_TO_UV
- GO_G1_DNA_DAMAGE_CHECKPOINT, GO_G1_DNA_DAMAGE_CHECKPOINT
- GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX, GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX
- GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
- GO_VIRION_ASSEMBLY, GO_VIRION_ASSEMBLY
- GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY, GO_UBIQUITIN_LIKE_PROTEIN_CONJUGATING_ENZYME_ACTIVITY