

GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION

- GO_HISTONE_UBIQUITINATION, GO_HISTONE_UBIQUITINATION
- GO_PROTEIN_MONOUBIQUITINATION, GO_PROTEIN_MONOUBIQUITINATION
- GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION_3_TO_LESION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION_3_TO_LESION
- GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
- GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING
- GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
- GO_TRANSLESION_SYNTHESIS, GO_TRANSLESION_SYNTHESIS
- GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DUPLEX_UNWINDING
- GO_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY, GO_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RE
- GO_ERROR_PRONE_TRANSLESION_SYNTHESIS, GO_ERROR_PRONE_TRANSLESION_SYNTHESIS
- GO_HISTONE_MONOUBIQUITINATION, GO_HISTONE_MONOUBIQUITINATION
- GO_DNA_POLYMERASE_ACTIVITY, GO_DNA_POLYMERASE_ACTIVITY
- GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_DAMAGE_RECOGNITION
- GO_HISTONE_H2A_UBIQUITINATION, GO_HISTONE_H2A_UBIQUITINATION
- GO_PYRIMIDINE_NUCLEOTIDE_METABOLIC_PROCESS, GO_PYRIMIDINE_NUCLEOTIDE_METABOLIC_PROCESS
- GO_CUL2_RING_UBIQUITIN_LIGASE_COMPLEX, GO_CUL2_RING_UBIQUITIN_LIGASE_COMPLEX
- GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION, GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION
- GO_HAIR_FOLLICLE_MATURATION, GO_HAIR_FOLLICLE_MATURATION
- GO_HAIR_CELL_DIFFERENTIATION, GO_HAIR_CELL_DIFFERENTIATION
- GO_REGULATION_OF_TELOMERE_CAPPING, GO_REGULATION_OF_TELOMERE_CAPPING
- GO_PYRIMIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_PYRIMIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
- GO_MITOTIC_RECOMBINATION, GO_MITOTIC_RECOMBINATION
- GO_PEROXISOMAL_TRANSPORT, GO_PEROXISOMAL_TRANSPORT