

ICASE_ACTIVITY, GO_HELICASE_ACTIVITY

- GO_HISTONE_BINDING, GO_HISTONE_BINDING
- GO_CHROMATIN_REMODELING, GO_CHROMATIN_REMODELING
- GO_DNA_GEOMETRIC_CHANGE, GO_DNA_GEOMETRIC_CHANGE
- GO_PROTEIN_DNA_COMPLEX, GO_PROTEIN_DNA_COMPLEX
- GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY
- GO_DNA_HELICASE_ACTIVITY, GO_DNA_HELICASE_ACTIVITY
- GO_NON_RECOMBINATIONAL_REPAIR, GO_NON_RECOMBINATIONAL_REPAIR
- GO_SINGLE_STRANDED_DNA_BINDING, GO_SINGLE_STRANDED_DNA_BINDING
- GO_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR, GO_TRANSCRIPTION_COUPLED_NUCLEOTIDE_EXCISION_REPAIR
- GO_NUCLEOTIDE_EXCISION_REPAIR, GO_NUCLEOTIDE_EXCISION_REPAIR
- GO_REPLICATION_FORK, GO_REPLICATION_FORK
- GO_ATP_DEPENDENT_CHROMATIN_REMODELING, GO_ATP_DEPENDENT_CHROMATIN_REMODELING
- GO_DAMAGED_DNA_BINDING, GO_DAMAGED_DNA_BINDING
- GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
- GO_ATPASE_COMPLEX, GO_ATPASE_COMPLEX
- GO_DNA_DAMAGE_RESPONSE_DETECTION_OF_DNA_DAMAGE, GO_DNA_DAMAGE_RESPONSE_DETECTION_OF_DNA_DAMAGE
- GO_RESPONSE_TO_IONIZING_RADIATION, GO_RESPONSE_TO_IONIZING_RADIATION
- GO_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GO_TRANSCRIPTION_BY_RNA_POLYMERASE_I
- GO_INTERSTRAND_CROSS_LINK_REPAIR, GO_INTERSTRAND_CROSS_LINK_REPAIR
- GO_NUCLEAR_REPLICATION_FORK, GO_NUCLEAR_REPLICATION_FORK
- GO_REGULATION_OF_DNA_RECOMBINATION, GO_REGULATION_OF_DNA_RECOMBINATION
- GO_TYPE_I_INTERFERON_PRODUCTION, GO_TYPE_I_INTERFERON_PRODUCTION
- GO_DOUBLE_STRANDED_RNA_BINDING, GO_DOUBLE_STRANDED_RNA_BINDING
- GO_DNA_REPAIR_COMPLEX, GO_DNA_REPAIR_COMPLEX
- GO_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX, GO_H4_HISTONE_ACETYLTRANSFERASE_COMPLEX
- GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
- GO_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION, GO_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION
- GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION, GO_POSITIVE_REGULATION_OF_VIRAL_TRANSCRIPTION
- GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_I_PROMOTER, GO_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_I_PROMOTER
- GO_DNA_STRAND_ELONGATION, GO_DNA_STRAND_ELONGATION
- GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION, GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION
- GO_METHYLATED_HISTONE_BINDING, GO_METHYLATED_HISTONE_BINDING
- GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS, GO_REGULATION_OF_DEFENSE_RESPONSE_TO_VIRUS
- GO_NEGATIVE_REGULATION_OF_DNA_REPAIR, GO_NEGATIVE_REGULATION_OF_DNA_REPAIR
- GO_HISTONE_DEACETYLASE_COMPLEX, GO_HISTONE_DEACETYLASE_COMPLEX
- GO_TELOMERASE_RNA_LOCALIZATION, GO_TELOMERASE_RNA_LOCALIZATION
- GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX, GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX
- GO_POSITIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_POSITIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
- GO_POSTREPLICATION_REPAIR, GO_POSTREPLICATION_REPAIR
- GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_BY_RNA_POLYMERASE_I
- GO_REGULATION_OF_VIRAL_TRANSCRIPTION, GO_REGULATION_OF_VIRAL_TRANSCRIPTION
- GO_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION, GO_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION
- GO_3_5_DNA_HELICASE_ACTIVITY, GO_3_5_DNA_HELICASE_ACTIVITY
- GO_TELOMERE_MAINTENANCE_VIA_SEMI_CONSERVATIVE_REPLICATION, GO_TELOMERE_MAINTENANCE_VIA_SEMI_CONSERVATIVE_REPLICATION
- GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
- GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
- GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION, GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION
- GO_DNA_DOUBLE_STRAND_BREAK_PROCESSING, GO_DNA_DOUBLE_STRAND_BREAK_PROCESSING
- GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_NEGATIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
- GO_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GO_NEGATIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
- GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY, GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
- GO_SWI_SNF_COMPLEX, GO_SWI_SNF_COMPLEX
- GO_NEGATIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GO_NEGATIVE_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
- GO_POSITIVE_REGULATION_OF_TELOMERASE_RNA_LOCALIZATION_TO_CAJAL_BODY, GO_POSITIVE_REGULATION_OF_TELOMERASE_RNA_LOCALIZATION_TO_CAJAL_BODY
- GO_MISMATCH_REPAIR_COMPLEX_BINDING, GO_MISMATCH_REPAIR_COMPLEX_BINDING
- GO_TELOMERIC_LOOP_DISASSEMBLY, GO_TELOMERIC_LOOP_DISASSEMBLY
- GO_FOUR_WAY_JUNCTION_HELICASE_ACTIVITY, GO_FOUR_WAY_JUNCTION_HELICASE_ACTIVITY
- GO_NONHOMOLOGOUS_END_JOINING_COMPLEX, GO_NONHOMOLOGOUS_END_JOINING_COMPLEX
- GO_HISTONE_H4_ACETYLTATION, GO_HISTONE_H4_ACETYLTATION
- GO_REPLISOME, GO_REPLISOME
- GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR, GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR
- GO_HISTONE_H2A_ACETYLTATION, GO_HISTONE_H2A_ACETYLTATION
- GO_FORMATION_OF_EXTRACHROMOSOMAL_CIRCULAR_DNA, GO_FORMATION_OF_EXTRACHROMOSOMAL_CIRCULAR_DNA
- GO_MLL1_2_COMPLEX, GO_MLL1_2_COMPLEX
- GO_INTERFERON_BETA_PRODUCTION, GO_INTERFERON_BETA_PRODUCTION
- GO_VIRAL_LATENCY, GO_VIRAL_LATENCY
- GO_U5_SNRNP, GO_U5_SNRNP
- GO_DNA_REPLICATION_ORIGIN_BINDING, GO_DNA_REPLICATION_ORIGIN_BINDING
- GO_NPBAF_COMPLEX, GO_NPBAF_COMPLEX
- GO_DNA_REPLICATION_INITIATION, GO_DNA_REPLICATION_INITIATION
- GO_RESPONSE_TO_DSRNA, GO_RESPONSE_TO_DSRNA
- GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY_IN_RESPONSE_TO_VIRUS, GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY_IN_RESPONSE_TO_VIRUS
- GO_TRANSPOSITION, GO_TRANSPOSITION
- GO_TRANSLESION_SYNTHESIS, GO_TRANSLESION_SYNTHESIS
- GO_TFIID_CLASS_TRANSCRIPTION_FACTOR_COMPLEX_BINDING, GO_TFIID_CLASS_TRANSCRIPTION_FACTOR_COMPLEX_BINDING
- GO_MULTI_ORGANISM_CELLULAR_PROCESS, GO_MULTI_ORGANISM_CELLULAR_PROCESS
- GO_H4_H2A_HISTONE_ACETYLTRANSFERASE_COMPLEX, GO_H4_H2A_HISTONE_ACETYLTRANSFERASE_COMPLEX