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GO_DNA_HELICASE_ACTIVITY, GO_DNA_HELICASE_ACTIVITY
GO_CELL_CYCLE_DNA_REPLICATION, GO_CELL_CYCLE_DNA_REPLICATION
GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_NUCLEOTIDE_EXCISION_REPAIR, GO_NUCLEOTIDE_EXCISION_REPAIR
GO_DNA_BIOSYNTHETIC_PROCESS, GO_DNA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_TELOMERE_MAINTENANCE, GO_REGULATION_OF_TELOMERE_MAINTENANCE
GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT
GO_CHROMATIN_SILENCING_AT_RDNA, GO_CHROMATIN_SILENCING_AT_RDNA
GO_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION, GO_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE
GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS, GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS
GO INTERSTRAND CROSS LINK REPAIR, GO INTERSTRAND CROSS LINK REPAIR
GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO TELOMERE MAINTENANCE VIA SEMI CONSERVATIVE REPLICATION, GO TELOMERE MAINTENANCE VIA SEMI CONSERVATIVE REPLICATION
GO_RESPONSE_TO_IONIZING_RADIATION, GO_RESPONSE_TO_IONIZING_RADIATION
GO_DAMAGED_DNA_BINDING, GO_DAMAGED_DNA_BINDING
GO_NON_RECOMBINATIONAL_REPAIR, GO_NON_RECOMBINATIONAL_REPAIR
GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
GO_DNA_REPLICATION_INITIATION, GO_DNA_REPLICATION_INITIATION
GO_POSTREPLICATION_REPAIR, GO_POSTREPLICATION_REPAIR
GO_REGULATION_OF_DNA_REPAIR, GO_REGULATION_OF_DNA_REPAIR
GO_DNA_STRAND_ELONGATION, GO_DNA_STRAND_ELONGATION
GO_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION, GO_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION
GO_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_ASSEMBLY, GO_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_ASSEMBLY
GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_SITE_OF_DOUBLE_STRAND_BREAK, GO_SITE_OF_DOUBLE_STRAND_BREAK
GO REGULATION OF DNA BIOSYNTHETIC PROCESS, GO REGULATION OF DNA BIOSYNTHETIC PROCESS
GO REGULATION OF DNA DEPENDENT DNA REPLICATION, GO REGULATION OF DNA DEPENDENT DNA REPLICATION
GO_NUCLEAR_NUCLEOSOME, GO_NUCLEAR_NUCLEOSOME
GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION, GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION
GO CHAPERONE COMPLEX, GO CHAPERONE COMPLEX
GO_TRANSLESION_SYNTHESIS, GO_TRANSLESION_SYNTHESIS
GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION,
GO_REGULATION_OF_TRANSCRIPTION_INVOLVED_IN_G1_S_TRANSITION_OF_MITOTIC_CELL_CYCLE, GO_REGULATION_OF_TRANSCRIPTION_INVOLVED_IN_G1_S_TRANSITION_OF_MITOTIC
GO_TELOMERE_CAPPING, GO_TELOMERE_CAPPING
GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR, GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR
GO FOUR WAY JUNCTION DNA BINDING, GO FOUR WAY JUNCTION DNA BINDING
GO_PROTEIN_C_TERMINUS_BINDING, GO_PROTEIN_C_TERMINUS_BINDING
GO_P53_BINDING, GO_P53_BINDING
GO_POSITIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION, GO_POSITIVE_REGULATION_OF_TYPE_I_INTERFERON_PRODUCTION
GO_PROTEIN_HETEROTETRAMERIZATION, GO_PROTEIN_HETEROTETRAMERIZATION
GO_ESTROGEN_RECEPTOR_BINDING, GO_ESTROGEN_RECEPTOR_BINDING
GO_G1_DNA_DAMAGE_CHECKPOINT, GO_G1_DNA_DAMAGE_CHECKPOINT
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
GO_HISTONE_DEACETYLASE_BINDING, GO_HISTONE_DEACETYLASE_BINDING
GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GO_DNA_POLYMERASE_COMPLEX, GO_DNA_POLYMERASE_COMPLEX
GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY, GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY
GO_REGULATION_OF_STEM_CELL_PROLIFERATION, GO_REGULATION_OF_STEM_CELL_PROLIFERATION
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
GO_MITOTIC_G2_M_TRANSITION_CHECKPOINT, GO_MITOTIC_G2_M_TRANSITION_CHECKPOINT
GO_3_5_DNA_HELICASE_ACTIVITY, GO_3_5_DNA_HELICASE_ACTIVITY
GO_DNA_REPAIR_COMPLEX, GO_DNA_REPAIR_COMPLEX
GO_REGULATION_OF_TELOMERE_CAPPING, GO_REGULATION_OF_TELOMERE_CAPPING
GO_BASE_EXCISION_REPAIR, GO_BASE_EXCISION_REPAIR
GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE
GO_PROTEIN_STABILIZATION, GO_PROTEIN_STABILIZATION
GO_DNA_SECONDARY_STRUCTURE_BINDING, GO_DNA_SECONDARY_STRUCTURE_BINDING
GO_RESPONSE_TO_GAMMA_RADIATION, GO_RESPONSE_TO_GAMMA_RADIATION
GO_MEGAKARYOCYTE_DIFFERENTIATION, GO_MEGAKARYOCYTE_DIFFERENTIATION
GO_SIGNAL_TRANSDUCTION_INVOLVED_IN_CELL_CYCLE_CHECKPOINT, GO_SIGNAL_TRANSDUCTION_INVOLVED_IN_CELL_CYCLE_CHECKPOINT
GO RESPONSE TO X RAY, GO RESPONSE TO X RAY
GO_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX, GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX
GO_DNA_REPLICATION_CHECKPOINT, GO_DNA_REPLICATION_CHECKPOINT
GO_TELOMERASE_RNA_LOCALIZATION, GO_TELOMERASE_RNA_LOCALIZATION
GO UNFOLDED PROTEIN BINDING, GO UNFOLDED PROTEIN BINDING
GO POSITIVE REGULATION OF RESPONSE TO DNA DAMAGE STIMULUS, GO POSITIVE REGULATION OF RESPONSE TO DNA DAMAGE STIMULUS
GO_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX, GO_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_ARREST
GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION, GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION
GO_3_UTR_MEDIATED_MRNA_DESTABILIZATION, GO_3_UTR_MEDIATED_MRNA_DESTABILIZATION
GO DEOXYRIBONUCLEASE ACTIVITY, GO DEOXYRIBONUCLEASE ACTIVITY
GO_MITOTIC_DNA_REPLICATION_CHECKPOINT, GO_MITOTIC_DNA_REPLICATION_CHECKPOINT
GO_PROTEIN_HETEROOLIGOMERIZATION, GO_PROTEIN_HETEROOLIGOMERIZATION
GO_PROTEIN_TETRAMERIZATION, GO_PROTEIN_TETRAMERIZATION
GO NEGATIVE REGULATION OF DNA METABOLIC PROCESS, GO NEGATIVE REGULATION OF DNA METABOLIC PROCESS
GO_NEGATIVE_REGULATION_OF_DNA_REPLICATION, GO_NEGATIVE_REGULATION_OF_DNA_REPLICATION
GO_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GO_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT
GO SYNAPSIS, GO SYNAPSIS
GO_MULTICELLULAR_ORGANISM_AGING, GO_MULTICELLULAR_ORGANISM_AGING
GO_MITOTIC_DNA_REPLICATION, GO_MITOTIC_DNA_REPLICATION
GO G2 DNA DAMAGE CHECKPOINT, GO G2 DNA DAMAGE CHECKPOINT
GO_CELLULAR_RESPONSE_TO_X_RAY, GO_CELLULAR_RESPONSE_TO_X_RAY
GO_MISMATCH_REPAIR_COMPLEX_BINDING, GO_MISMATCH_REPAIR_COMPLEX_BINDING
GO_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
GO_TELOMERE_MAINTENANCE_VIA_RECOMBINATION, GO_TELOMERE_MAINTENANCE_VIA_RECOMBINATION
GO_REGULATION_OF_METANEPHROS_DEVELOPMENT, GO_REGULATION_OF_METANEPHROS_DEVELOPMENT
GO MALE MEIOSIS I, GO MALE MEIOSIS I
GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
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GO DNA DEPENDENT DNA REPLICATION, GO DNA DEPENDENT DNA REPLICATION

GO_DNA_DEPENDENT_ATPASE_ACTIVITY, GO_DNA_DEPENDENT_ATPASE_ACTIVITY

GO_SINGLE_STRANDED_DNA_BINDING, GO_SINGLE_STRANDED_DNA_BINDING

GO_DNA_PACKAGING_COMPLEX, GO_DNA_PACKAGING_COMPLEX

GO_RNA_POLYMERASE_COMPLEX, GO_RNA_POLYMERASE_COMPLEX

GO_RECOMBINATIONAL_REPAIR, GO_RECOMBINATIONAL_REPAIR

GO DNA INTEGRITY CHECKPOINT, GO DNA INTEGRITY CHECKPOINT

GO_DNA_GEOMETRIC_CHANGE, GO_DNA_GEOMETRIC_CHANGE

GO_PROTEIN_DNA_COMPLEX, GO_PROTEIN_DNA_COMPLEX

GO_REPLICATION_FORK, GO_REPLICATION_FORK

ORGANIZATION, GO TELOMERE ORGANIZATION