

ORGANIZATION, GO_TELOMERE_ORGANIZATION

GO_SISTER_CHROMATID_SEGREGATION, GO_SISTER_CHROMATID_SEGREGATION
GO_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_MITOTIC_SISTER_CHROMATID_SEGREGATION
GO_DNA_DEPENDENT_DNA_REPLICATION, GO_DNA_DEPENDENT_DNA_REPLICATION
GO_RECOMBINATIONAL_REPAIR, GO_RECOMBINATIONAL_REPAIR
GO_NUCLEOTIDE_EXCISION_REPAIR, GO_NUCLEOTIDE_EXCISION_REPAIR
GO_DNA_DEPENDENT_ATPASE_ACTIVITY, GO_DNA_DEPENDENT_ATPASE_ACTIVITY
GO_CHROMOSOME_TELOMERIC_REGION, GO_CHROMOSOME_TELOMERIC_REGION
GO_DNA_GEOMETRIC_CHANGE, GO_DNA_GEOMETRIC_CHANGE
GO_NON_RECOMBINATIONAL_REPAIR, GO_NON_RECOMBINATIONAL_REPAIR
GO_DAMAGED_DNA_BINDING, GO_DAMAGED_DNA_BINDING
GO_DNA_BIOSYNTHETIC_PROCESS, GO_DNA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_DNA_HELICASE_ACTIVITY, GO_DNA_HELICASE_ACTIVITY
GO_REGULATION_OF_TELOMERE_MAINTENANCE, GO_REGULATION_OF_TELOMERE_MAINTENANCE
GO_SINGLE_STRANDED_DNA_BINDING, GO_SINGLE_STRANDED_DNA_BINDING
GO_SITE_OF_DNA_DAMAGE, GO_SITE_OF_DNA_DAMAGE
GO_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_REGULATION_OF_GENE_SILENCING, GO_REGULATION_OF_GENE_SILENCING
GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION, GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION
GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
GO_PROTEIN_DNA_COMPLEX, GO_PROTEIN_DNA_COMPLEX
GO_RNA_POLYMERASE_COMPLEX, GO_RNA_POLYMERASE_COMPLEX
GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION, GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION
GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS, GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS
GO_CELL_CYCLE_DNA_REPLICATION, GO_CELL_CYCLE_DNA_REPLICATION
GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
GO_DNA_INTEGRITY_CHECKPOINT, GO_DNA_INTEGRITY_CHECKPOINT
GO_INTERSTRAND_CROSS_LINK_REPAIR, GO_INTERSTRAND_CROSS_LINK_REPAIR
GO_MITOTIC_CELL_CYCLE_CHECKPOINT, GO_MITOTIC_CELL_CYCLE_CHECKPOINT
GO_REPLICATION_FORK, GO_REPLICATION_FORK
GO_DNA_PACKAGING_COMPLEX, GO_DNA_PACKAGING_COMPLEX
GO_CHROMATIN_REMODELING, GO_CHROMATIN_REMODELING
GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
GO_HETEROCHROMATIN, GO_HETEROCHROMATIN
GO_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION, GO_DNA_REPLICATION_DEPENDENT_NUCLEOSOME_ORGANIZATION
GO_EXONUCLEASE_ACTIVITY, GO_EXONUCLEASE_ACTIVITY
GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_CHROMATIN_ORGANIZATION_INVOLVED_IN_REGULATION_OF_TRANSCRIPTION, GO_CHROMATIN_ORGANIZATION_INVOLVED_IN_REGULATION_OF_TRANSCRIPTION
GO_NUCLEASE_ACTIVITY, GO_NUCLEASE_ACTIVITY
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_GAP_FILLING
GO_NUCLEOSOME_ORGANIZATION, GO_NUCLEOSOME_ORGANIZATION
GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR, GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR
GO_DNA_REPLICATION_INITIATION, GO_DNA_REPLICATION_INITIATION
GO_CHROMOSOME_SEPARATION, GO_CHROMOSOME_SEPARATION
GO_TELOMERE_MAINTENANCE_VIA_SEMI_CONSERVATIVE_REPLICATION, GO_TELOMERE_MAINTENANCE_VIA_SEMI_CONSERVATIVE_REPLICATION
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_POSTREPLICATION_REPAIR, GO_POSTREPLICATION_REPAIR
GO_SITE_OF_DOUBLE_STRAND_BREAK, GO_SITE_OF_DOUBLE_STRAND_BREAK
GO_CHROMATIN_ASSEMBLY, GO_CHROMATIN_ASSEMBLY
GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY, GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY
GO_SNORNA_BINDING, GO_SNORNA_BINDING
GO_REGULATION_OF_DNA_REPAIR, GO_REGULATION_OF_DNA_REPAIR
GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PRIME_TO_3_PRIME_DIRECTIONALITY, GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PRIME_TO_3_PRIME_DIRECTIONALITY
GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
GO_DNA_REPAIR_COMPLEX, GO_DNA_REPAIR_COMPLEX
GO_TRANSLESION_SYNTHESIS, GO_TRANSLESION_SYNTHESIS
GO_MISMATCH_REPAIR, GO_MISMATCH_REPAIR
GO_RESPONSE_TO_IONIZING_RADIATION, GO_RESPONSE_TO_IONIZING_RADIATION
GO_NUCLEAR_HETEROCHROMATIN, GO_NUCLEAR_HETEROCHROMATIN
GO_SISTER_CHROMATID_COHESION, GO_SISTER_CHROMATID_COHESION
GO_CHROMATIN_SILENCING_AT_RDNA, GO_CHROMATIN_SILENCING_AT_RDNA
GO_METAL_CLUSTER_BINDING, GO_METAL_CLUSTER_BINDING
GO_DNA_STRAND_ELONGATION, GO_DNA_STRAND_ELONGATION
GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
GO_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION, GO_DNA_STRAND_ELONGATION_INVOLVED_IN_DNA_REPLICATION
GO_4_IRON_4_SULFUR_CLUSTER_BINDING, GO_4_IRON_4_SULFUR_CLUSTER_BINDING
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING
GO_BASE_EXCISION_REPAIR, GO_BASE_EXCISION_REPAIR
GO_3_5_DNA_HELICASE_ACTIVITY, GO_3_5_DNA_HELICASE_ACTIVITY
GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY, GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY
GO_PROTEIN_HETEROTETRAMERIZATION, GO_PROTEIN_HETEROTETRAMERIZATION
GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY
GO_PROTEIN_TETRAMERIZATION, GO_PROTEIN_TETRAMERIZATION
GO_3_5_EXONUCLEASE_ACTIVITY, GO_3_5_EXONUCLEASE_ACTIVITY
GO_PROTEIN_C_TERMINUS_BINDING, GO_PROTEIN_C_TERMINUS_BINDING
GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_REGULATION_OF_TELOMERASE_ACTIVITY, GO_REGULATION_OF_TELOMERASE_ACTIVITY
GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION
GO_NUCLEOSOMAL_DNA_BINDING, GO_NUCLEOSOMAL_DNA_BINDING
GO_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GO_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_ASSEMBLY, GO_NUCLEOTIDE_EXCISION_REPAIR_PREINCISION_COMPLEX_ASSEMBLY
GO_DNA_POLYMERASE_BINDING, GO_DNA_POLYMERASE_BINDING
GO_PROTEIN_LOCALIZATION_TO_NUCLEOLUS, GO_PROTEIN_LOCALIZATION_TO_NUCLEOLUS
GO_NEGATIVE_REGULATION_OF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GO_NEGATIVE_REGULATION_OF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
GO_REGULATION_OF_SISTER_CHROMATID_COHESION, GO_REGULATION_OF_SISTER_CHROMATID_COHESION
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
GO_ATP_DEPENDENT_CHROMATIN_REMODELING, GO_ATP_DEPENDENT_CHROMATIN_REMODELING
GO_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY, GO_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY
GO_PML_BODY, GO_PML_BODY
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_DNA_REPLICATION_INDEPENDENT_NUCLEOSOME_ORGANIZATION, GO_DNA_REPLICATION_INDEPENDENT_NUCLEOSOME_ORGANIZATION
GO_POSITIVE_REGULATION_OF_DNA_REPAIR, GO_POSITIVE_REGULATION_OF_DNA_REPAIR
GO_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
GO_NUCLEAR_NUCLEOSOME, GO_NUCLEAR_NUCLEOSOME
GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY, GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY
GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION, GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION
GO_MANGANESE_ION_BINDING, GO_MANGANESE_ION_BINDING
GO_RNA_SURVEILLANCE, GO_RNA_SURVEILLANCE
GO_REGULATION_OF_DNA_RECOMBINATION, GO_REGULATION_OF_DNA_RECOMBINATION
GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_NUCLEAR_DIVISION, GO_REGULATION_OF_NUCLEAR_DIVISION
GO_STEROID_HORMONE_RECEPTOR_BINDING, GO_STEROID_HORMONE_RECEPTOR_BINDING
GO_CHROMATIN_REMODELING_AT_CENTROMERE, GO_CHROMATIN_REMODELING_AT_CENTROMERE
GO_TELOMERE_CAPPING, GO_TELOMERE_CAPPING
GO_REPLICATION_FORK_PROCESSING, GO_REPLICATION_FORK_PROCESSING
GO_TYPE_I_INTERFERON_PRODUCTION, GO_TYPE_I_INTERFERON_PRODUCTION
GO_CENTROMERE_COMPLEX_ASSEMBLY, GO_CENTROMERE_COMPLEX_ASSEMBLY
GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS, GO_SOMATIC_DIVERSIFICATION_OF_IMMUNE_RECEPTORS
GO_MITOTIC_SISTER_CHROMATID_COHESION, GO_MITOTIC_SISTER_CHROMATID_COHESION
GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX, GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX
GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
GO_NUCLEAR_RECEPTOR_BINDING, GO_NUCLEAR_RECEPTOR_BINDING
GO_DEOXYRIBONUCLEASE_ACTIVITY, GO_DEOXYRIBONUCLEASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
GO_RESPONSE_TO_UV, GO_RESPONSE_TO_UV
GO_CELLULAR_RESPONSE_TO_HEAT, GO_CELLULAR_RESPONSE_TO_HEAT
GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION, GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
GO_CELL_RECOGNITION, GO_CELL_RECOGNITION
GO_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
GO_REGULATION_OF_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_REGULATION_OF_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_DNA_POLYMERASE_COMPLEX, GO_DNA_POLYMERASE_COMPLEX
GO_RIBONUCLEASE_ACTIVITY, GO_RIBONUCLEASE_ACTIVITY
GO_FOUR_WAY_JUNCTION_DNA_BINDING, GO_FOUR_WAY_JUNCTION_DNA_BINDING
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION_3_TO_LESION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION_3_TO_LESION
GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS, GO_POSITIVE_REGULATION_OF_PROTEIN_LOCALIZATION_TO_NUCLEUS
GO_PROTEIN_DESTABILIZATION, GO_PROTEIN_DESTABILIZATION
GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT, GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT
GO_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION, GO_REGULATION_OF_NUCLEAR_CELL_CYCLE_DNA_REPLICATION
GO_PROTEIN_ADP_RIBOSYLATION, GO_PROTEIN_ADP_RIBOSYLATION
GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT