

ORGANIZATION, GO_TELOMERE_ORGANIZATION

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_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_SISTER_CHROMATID_SEGREGATION, GO_SISTER_CHROMATID_SEGREGATION
GO_CHROMOSOME_TELOMERIC_REGION, GO_CHROMOSOME_TELOMERIC_REGION
GO_DNA_BIOSYNTHETIC_PROCESS, GO_DNA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_DNA_REPAIR, GO_REGULATION_OF_DNA_REPAIR
GO_NON_RECOMBINATIONAL_REPAIR, GO_NON_RECOMBINATIONAL_REPAIR
GO_NUCLEOTIDE_EXCISION_REPAIR, GO_NUCLEOTIDE_EXCISION_REPAIR
GO_DNA_DEPENDENT_ATPASE_ACTIVITY, GO_DNA_DEPENDENT_ATPASE_ACTIVITY
GO_RNA_POLYMERASE_COMPLEX, GO_RNA_POLYMERASE_COMPLEX
GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY, GO_NUCLEOTIDYLTRANSFERASE_ACTIVITY
GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_DNA_INTEGRITY_CHECKPOINT, GO_DNA_INTEGRITY_CHECKPOINT
GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
GO_RESPONSE_TO_IONIZING_RADIATION, GO_RESPONSE_TO_IONIZING_RADIATION
GO_REGULATION_OF_GENE_SILENCING, GO_REGULATION_OF_GENE_SILENCING
GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION, GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION
GO_DNA_HELICASE_ACTIVITY, GO_DNA_HELICASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT, GO_REGULATION_OF_CELLULAR_RESPONSE_TO_HEAT
GO_SITE_OF_DNA_DAMAGE, GO_SITE_OF_DNA_DAMAGE
GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS, GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_DAMAGED_DNA_BINDING, GO_DAMAGED_DNA_BINDING
GO_CELL_CYCLE_DNA_REPLICATION, GO_CELL_CYCLE_DNA_REPLICATION
GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
GO_DNA_GEOMETRIC_CHANGE, GO_DNA_GEOMETRIC_CHANGE
GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION, GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION
GO_REGULATION_OF_TELOMERE_MAINTENANCE, GO_REGULATION_OF_TELOMERE_MAINTENANCE
GO_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_SINGLE_STRANDED_DNA_BINDING, GO_SINGLE_STRANDED_DNA_BINDING
GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION, GO_CELLULAR_RESPONSE_TO_IONIZING_RADIATION
GO_INTERSTRAND_CROSS_LINK_REPAIR, GO_INTERSTRAND_CROSS_LINK_REPAIR
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
GO_REPLICATION_FORK, GO_REPLICATION_FORK
GO_MITOTIC_CELL_CYCLE_CHECKPOINT, GO_MITOTIC_CELL_CYCLE_CHECKPOINT
GO_DNA_REPLICATION_INITIATION, GO_DNA_REPLICATION_INITIATION
GO_EXONUCLEASE_ACTIVITY, GO_EXONUCLEASE_ACTIVITY
GO_REGULATION_OF_DNA_REPLICATION, GO_REGULATION_OF_DNA_REPLICATION
GO_NUCLEASE_ACTIVITY, GO_NUCLEASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
GO_HETEROCHROMATIN, GO_HETEROCHROMATIN
GO_CHROMATIN_ORGANIZATION_INVOLVED_IN_REGULATION_OF_TRANSCRIPTION, GO_CHROMATIN_ORGANIZATION_INVOLVED_IN_REGULATION_OF_TRANSCRIPTION
GO_CHROMOSOME_SEPARATION, GO_CHROMOSOME_SEPARATION
GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
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_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_EXONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_P
GO_CHROMATIN_REMODELING, GO_CHROMATIN_REMODELING
GO_REGULATION_OF_NUCLEAR_DIVISION, GO_REGULATION_OF_NUCLEAR_DIVISION
GO_POSITIVE_REGULATION_OF_DNA_REPAIR, GO_POSITIVE_REGULATION_OF_DNA_REPAIR
GO_POSTREPLICATION_REPAIR, GO_POSTREPLICATION_REPAIR
GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
GO_NUCLEOSOME_ORGANIZATION, GO_NUCLEOSOME_ORGANIZATION
GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
GO_REGULATION_OF_SISTER_CHROMATID_COHESION, GO_REGULATION_OF_SISTER_CHROMATID_COHESION
GO_REGULATION_OF_CHROMOSOME_SEPARATION, GO_REGULATION_OF_CHROMOSOME_SEPARATION
GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY
GO_DEOXYRIBONUCLEASE_ACTIVITY, GO_DEOXYRIBONUCLEASE_ACTIVITY
GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR, GO_DNA_SYNTHESIS_INVOLVED_IN_DNA_REPAIR
GO_DNA_PACKAGING_COMPLEX, GO_DNA_PACKAGING_COMPLEX
GO_ENDODEOXYRIBONUCLEASE_ACTIVITY, GO_ENDODEOXYRIBONUCLEASE_ACTIVITY
GO_PROTEIN_DNA_COMPLEX, GO_PROTEIN_DNA_COMPLEX
GO_CELLULAR_RESPONSE_TO_HEAT, GO_CELLULAR_RESPONSE_TO_HEAT
GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT, GO_REGULATION_OF_CELL_CYCLE_CHECKPOINT
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
GO_3_5_EXONUCLEASE_ACTIVITY, GO_3_5_EXONUCLEASE_ACTIVITY
GO_PROTEIN_C_TERMINUS_BINDING, GO_PROTEIN_C_TERMINUS_BINDING
GO_CHROMATIN_ASSEMBLY, GO_CHROMATIN_ASSEMBLY
GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION, GO_DNA_UNWINDING_INVOLVED_IN_DNA_REPLICATION
GO_SITE_OF_DOUBLE_STRAND_BREAK, GO_SITE_OF_DOUBLE_STRAND_BREAK
GO_NUCLEAR_HETEROCHROMATIN, GO_NUCLEAR_HETEROCHROMATIN
GO_TRANSLESION_SYNTHESIS, GO_TRANSLESION_SYNTHESIS
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE
GO_CELLULAR_RESPONSE_TO_RADIATION, GO_CELLULAR_RESPONSE_TO_RADIATION
GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_P
GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY, GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY
GO_SISTER_CHROMATID_COHESION, GO_SISTER_CHROMATID_COHESION
GO_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_REGULATION_OF_CHROMATIN_ORGANIZATION
GO_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GO_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
GO_CENTROMERE_COMPLEX_ASSEMBLY, GO_CENTROMERE_COMPLEX_ASSEMBLY
GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
GO_RESPONSE_TO_UV, GO_RESPONSE_TO_UV
GO_REPLICATION_FORK_PROCESSING, GO_REPLICATION_FORK_PROCESSING
GO_NEGATIVE_REGULATION_OF_NUCLEAR_DIVISION, GO_NEGATIVE_REGULATION_OF_NUCLEAR_DIVISION
GO_PROTEIN_ADP_RIBOSYLASE_ACTIVITY, GO_PROTEIN_ADP_RIBOSYLASE_ACTIVITY
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS
GO_CELL_RECOGNITION, GO_CELL_RECOGNITION
GO_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE, GO_TELOMERE_MAINTENANCE_IN_RESPONSE_TO_DNA_DAMAGE
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE
GO_SNORNA_BINDING, GO_SNORNA_BINDING
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_HOMOLOGOUS_RECOMBINATION
GO_PHOSPHATASE_BINDING, GO_PHOSPHATASE_BINDING
GO_MITOCHONDRIAL_GENOME_MAINTENANCE, GO_MITOCHONDRIAL_GENOME_MAINTENANCE
GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
GO_RESPONSE_TO_GAMMA_RADIATION, GO_RESPONSE_TO_GAMMA_RADIATION
GO_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY, GO_INTERLEUKIN_7_MEDIATED_SIGNALING_PATHWAY
GO_DNA_POLYMERASE_BINDING, GO_DNA_POLYMERASE_BINDING
GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC, GO_RNA_PHOSPHODIESTER_BOND_HYDROLYSIS_EXONUCLEOLYTIC
GO_HISTONE_DEACETYLASE_BINDING, GO_HISTONE_DEACETYLASE_BINDING
GO_CELLULAR_RESPONSE_TO_SALT_STRESS, GO_CELLULAR_RESPONSE_TO_SALT_STRESS
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_ENDONUCLEASE_ACTIVITY, GO_ENDONUCLEASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION
GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
GO_RESPONSE_TO_INTERLEUKIN_7, GO_RESPONSE_TO_INTERLEUKIN_7
GO_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT, GO_REGULATION_OF_EARLY_ENDOSOME_TO_LATE_ENDOSOME_TRANSPORT
GO_MEGAKARYOCYTE_DIFFERENTIATION, GO_MEGAKARYOCYTE_DIFFERENTIATION
GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT
GO_EXODEOXYRIBONUCLEASE_ACTIVITY, GO_EXODEOXYRIBONUCLEASE_ACTIVITY
GO_RESPONSE_TO_HEAT, GO_RESPONSE_TO_HEAT
GO_MITOTIC_SISTER_CHROMATID_COHESION, GO_MITOTIC_SISTER_CHROMATID_COHESION
GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR, GO_DNA_DAMAGE_RESPONSE_SIGNAL_TRANSDUCTION_BY_P53_CLASS_MEDIATOR
GO_EXORIBONUCLEASE_ACTIVITY, GO_EXORIBONUCLEASE_ACTIVITY
GO_PML_BODY, GO_PML_BODY
GO_ERROR_PRONE_TRANSLESION_SYNTHESIS, GO_ERROR_PRONE_TRANSLESION_SYNTHESIS
GO_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE, GO_PROTECTION_FROM_NON_HOMOLOGOUS_END_JOINING_AT_TELOMERE
GO_DNA_POLYMERASE_ACTIVITY, GO_DNA_POLYMERASE_ACTIVITY
GO_CENTRIOLE_CENTRIOLE_COHESION, GO_CENTRIOLE_CENTRIOLE_COHESION
GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE, GO_MAINTENANCE_OF_PROTEIN_LOCALIZATION_IN_ORGANELLE
GO_DNA_DOUBLE_STRAND_BREAK_PROCESSING, GO_DNA_DOUBLE_STRAND_BREAK_PROCESSING
GO_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX, GO_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX
GO_ERROR_FREE_TRANSLESION_SYNTHESIS, GO_ERROR_FREE_TRANSLESION_SYNTHESIS
GO_MEIOTIC_CELL_CYCLE_PROCESS, GO_MEIOTIC_CELL_CYCLE_PROCESS
GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX, GO_ADA2_GCN5_ADA3_TRANSCRIPTION_ACTIVATOR_COMPLEX
GO_DNA_REPLICATION_CHECKPOINT, GO_DNA_REPLICATION_CHECKPOINT
GO_ATP_DEPENDENT_CHROMATIN_REMODELING, GO_ATP_DEPENDENT_CHROMATIN_REMODELING
GO_REGULATION_OF_TELOMERE_CAPPING, GO_REGULATION_OF_TELOMERE_CAPPING
GO_STEROID_HORMONE_RECEPTOR_BINDING, GO_STEROID_HORMONE_RECEPTOR_BINDING
GO_DNA_SECONDARY_STRUCTURE_BINDING, GO_DNA_SECONDARY_STRUCTURE_BINDING
GO_ANNEALING_ACTIVITY, GO_ANNEALING_ACTIVITY
GO_REGULATION_OF_TRANSCRIPTION_INVOLVED_IN_G1_S_TRANSITION_OF_MITOTIC_CELL_CYCLE, GO_REGULATION_OF_TRANSCRIPTION_INVOLVED_IN_G1_S_TRANSITION_OF_MITOTIC_CELL_CYCLE
GO_POSITIVE_REGULATION_OF_TYPE_1_INTERFERON_PRODUCTION, GO_POSITIVE_REGULATION_OF_TYPE_1_INTERFERON_PRODUCTION
GO_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION
GO_BASE_EXCISION_REPAIR, GO_BASE_EXCISION_REPAIR
GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR, GO_GLOBAL_GENOME_NUCLEOTIDE_EXCISION_REPAIR
GO_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION, GO_REGULATION_OF_MEGAKARYOCYTE_DIFFERENTIATION
GO_NEGATIVE_REGULATION_OF_DNA_REPAIR, GO_NEGATIVE_REGULATION_OF_DNA_REPAIR
GO_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY, GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY
GO_REGULATION_OF_DNA_DAMAGE_CHECKPOINT, GO_REGULATION_OF_DNA_DAMAGE_CHECKPOINT
GO_POSITIVE_REGULATION_OF_BINDING, GO_POSITIVE_REGULATION_OF_BINDING
GO_FIBROBLAST_PROLIFERATION, GO_FIBROBLAST_PROLIFERATION
GO_MALE_MEIOTIC_NUCLEAR_DIVISION, GO_MALE_MEIOTIC_NUCLEAR_DIVISION
GO_P53_BINDING, GO_P53_BINDING
GO_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
GO_RESPONSE_TO_CADMIUM_ION, GO_RESPONSE_TO_CADMIUM_ION
GO_NADPLUS_ADP_RIBOSYLTRANSFERASE_ACTIVITY, GO_NADPLUS_ADP_RIBOSYLTRANSFERASE_ACTIVITY
GO_VIRAL_LATENCY, GO_VIRAL_LATENCY
GO_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION, GO_POSITIVE_REGULATION_OF_SISTER_CHROMATID_COHESION
GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY
GO_HISTONE_PHOSPHORYLATION, GO_HISTONE_PHOSPHORYLATION
GO_NCRNA_CATABOLIC_PROCESS, GO_NCRNA_CATABOLIC_PROCESS
GO_MAINTENANCE_OF_SISTER_CHROMATID_COHESION, GO_MAINTENANCE_OF_SISTER_CHROMATID_COHESION
GO_CELLULAR_RESPONSE_TO_X_RAY, GO_CELLULAR_RESPONSE_TO_X_RAY
GO_DNA_DIRECTED_DNA_POLYMERASE_ACTIVITY, GO_DNA_DIRECTED_DNA_POLYMERASE_ACTIVITY
GO_BINDING_OF_SPERM_TO_ZONA_PELLUCIDA, GO_BINDING_OF_SPERM_TO_ZONA_PELLUCIDA
GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION, GO_NUCLEOTIDE_EXCISION_REPAIR_DNA_INCISION
GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_REGULATION_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
GO_DNA_REPLICATION_INDEPENDENT_NUCLEOSOME_ORGANIZATION, GO_DNA_REPLICATION_INDEPENDENT_NUCLEOSOME_ORGANIZATION
GO_DNA_LIGATION, GO_DNA_LIGATION
GO_PROTEIN_STABILIZATION, GO_PROTEIN_STABILIZATION
GO_CHROMATIN_REMODELING_AT_CENTROMERE, GO_CHROMATIN_REMODELING_AT_CENTROMERE
GO_PROTEIN_AUTO_ADP_RIBOSYLATION, GO_PROTEIN_AUTO_ADP_RIBOSYLATION
GO_REGULATION_OF_DNA_RECOMBINATION, GO_REGULATION_OF_DNA_RECOMBINATION
GO_TELOMERE_CAPPING, GO_TELOMERE_CAPPING
GO_REGULATION_OF_CHROMATIN_BINDING, GO_REGULATION_OF_CHROMATIN_BINDING
GO_DNA_STRAND_ELONGATION, GO_DNA_STRAND_ELONGATION
GO_RNA_SURVEILLANCE, GO_RNA_SURVEILLANCE
GO_POSITIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION, GO_POSITIVE_REGULATION_OF_FIBROBLAST_PROLIFERATION
GO_CROSSOVER_JUNCTION_ENDODEOXYRIBONUCLEASE_ACTIVITY, GO_CROSSOVER_JUNCTION_ENDODEOXYRIBONUCLEASE_ACTIVITY
GO_MITOTIC_DNA_REPLICATION_CHECKPOINT, GO_MITOTIC_DNA_REPLICATION_CHECKPOINT
GO_MITOCHONDRIAL_DNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_DNA_METABOLIC_PROCESS
GO_REGULATION_OF_CELL_CYCLE_ARREST, GO_REGULATION_OF_CELL_CYCLE_ARREST
GO_TELOMERASE_RNA_LOCALIZATION, GO_TELOMERASE_RNA_LOCALIZATION
GO_HISTONE_EXCHANGE, GO_HISTONE_EXCHANGE
GO_FLAP_ENDONUCLEASE_ACTIVITY, GO_FLAP_ENDONUCLEASE_ACTIVITY
GO_FOUR_WAY_JUNCTION_DNA_BINDING, GO_FOUR_WAY_JUNCTION_DNA_BINDING
GO_CELLULAR_RESPONSE_TO_GAMMA_RADIATION, GO_CELLULAR_RESPONSE_TO_GAMMA_RADIATION
GO_RRNA_PSEUDOURIDINE_SYNTHESIS, GO_RRNA_PSEUDOURIDINE_SYNTHESIS
GO_POSITIVE_REGULATION_OF_TELOMERASE_RNA_LOCALIZATION_TO_CAJAL_BODY, GO_POSITIVE_REGULATION_OF_TELOMERASE_RNA_LOCALIZATION_TO_CAJAL_BODY
GO_NEGATIVE_REGULATION_OF_TELOMERASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_TELOMERASE_ACTIVITY
GO_TRACHEA_MORPHOGENESIS, GO_TRACHEA_MORPHOGENESIS
GO_SEX_CHROMOSOME, GO_SEX_CHROMOSOME
GO_DNA_MODIFICATION, GO_DNA_MODIFICATION