CHROMOSOME_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION

GO_RNA_3_END_PROCESSING, GO_RNA_3_END_PROCESSING
GO_TELOMERE_ORGANIZATION, GO_TELOMERE_ORGANIZATION
GO_MRNA_3_END_PROCESSING, GO_MRNA_3_END_PROCESSING
GO_ESTABLISHMENT_OF_RNA_LOCALIZATION, GO_ESTABLISHMENT_OF_RNA_LOCALIZATION
GO_CHROMOSOME_TELOMERIC_REGION, GO_CHROMOSOME_TELOMERIC_REGION
GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION, GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE
GO_REGULATION_OF_SISTER_CHROMATID_COHESION, GO_REGULATION_OF_SISTER_CHROMATID_COHESION
GO_MITOTIC_CELL_CYCLE_CHECKPOINT, GO_MITOTIC_CELL_CYCLE_CHECKPOINT
GO_REGULATION_OF_TELOMERE_MAINTENANCE, GO_REGULATION_OF_TELOMERE_MAINTENANCE
GO_REPLICATION_FORK, GO_REPLICATION_FORK
GO_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION, GO_NEGATIVE_REGULATION_OF_DNA_RECOMBINATION
GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE
GO_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_NUCLEAR_MATRIX, GO_NUCLEAR_MATRIX
GO_CHROMATIN_REMODELING, GO_CHROMATIN_REMODELING
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
GO_NUCLEAR_PERIPHERY, GO_NUCLEAR_PERIPHERY
GO_SISTER_CHROMATID_SEGREGATION, GO_SISTER_CHROMATID_SEGREGATION
GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
GO_RESPONSE_TO_GAMMA_RADIATION, GO_RESPONSE_TO_GAMMA_RADIATION
GO_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMATIN_ORGANIZATION
GO_NUCLEAR_EXPORT, GO_NUCLEAR_EXPORT
GO_HISTONE_DEACETYLASE_COMPLEX, GO_HISTONE_DEACETYLASE_COMPLEX
GO_SAGA_TYPE_COMPLEX, GO_SAGA_TYPE_COMPLEX
GO_PROMOTER_SPECIFIC_CHROMATIN_BINDING, GO_PROMOTER_SPECIFIC_CHROMATIN_BINDING
GO_RECOMBINATIONAL_REPAIR, GO_RECOMBINATIONAL_REPAIR
GO_MRNA_TRANSPORT, GO_MRNA_TRANSPORT
GO_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_MITOTIC_SISTER_CHROMATID_SEGREGATION
GO_PEPTIDYL_LYSINE_ACETYLATION, GO_PEPTIDYL_LYSINE_ACETYLATION
GO_POSITIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE, GO_POSITIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_TELOMERE
GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY, GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY
GO_PROTEIN_ACETYLTRANSFERASE_COMPLEX, GO_PROTEIN_ACETYLTRANSFERASE_COMPLEX
GO_TELOMERASE_HOLOENZYME_COMPLEX, GO_TELOMERASE_HOLOENZYME_COMPLEX
GO_SYNAPTIC_VESICLE_TRANSPORT, GO_SYNAPTIC_VESICLE_TRANSPORT
GO_DNA_BIOSYNTHETIC_PROCESS, GO_DNA_BIOSYNTHETIC_PROCESS
GO_CELL_CYCLE_DNA_REPLICATION, GO_CELL_CYCLE_DNA_REPLICATION
GO_SPINDLE_ORGANIZATION, GO_SPINDLE_ORGANIZATION
GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
GO_PROTEIN_SUMOYLATION, GO_PROTEIN_SUMOYLATION
GO_G1_DNA_DAMAGE_CHECKPOINT, GO_G1_DNA_DAMAGE_CHECKPOINT
GO_DNA_DEPENDENT_DNA_REPLICATION, GO_DNA_DEPENDENT_DNA_REPLICATION
GO_TELOMERASE_RNA_LOCALIZATION, GO_TELOMERASE_RNA_LOCALIZATION
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION, GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION
GO_SISTER_CHROMATID_COHESION, GO_SISTER_CHROMATID_COHESION
GO_PROTEIN_LOCALIZATION_TO_NUCLEOPLASM, GO_PROTEIN_LOCALIZATION_TO_NUCLEOPLASM
GO_REGULATION_OF_HELICASE_ACTIVITY, GO_REGULATION_OF_HELICASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RESPONSE_TO_CHEMICAL_STIMULUS, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_INVOLVED_IN_CELLULAR_RE
GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS, GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS
GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_REGULATION_OF_CHROMATIN_ORGANIZATION