GO_SISTER_CHROMATID_SEGREGATION, GO_SISTER_CHROMATID_SEGREGATION
GO_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_MITOTIC_SISTER_CHROMATID_SEGREGATION
GO_SPINDLE_POLE, GO_SPINDLE_POLE
GO_SPINDLE_ORGANIZATION, GO_SPINDLE_ORGANIZATION
GO_CILIARY_BASAL_BODY_PLASMA_MEMBRANE_DOCKING, GO_CILIARY_BASAL_BODY_PLASMA_MEMBRANE_DOCKING
GO_CHROMOSOME_TELOMERIC_REGION, GO_CHROMOSOME_TELOMERIC_REGION
GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING, GO_REGULATION_OF_POSTTRANSCRIPTIONAL_GENE_SILENCING
GO_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS, GO_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS
GO_KINETOCHORE, GO_KINETOCHORE
GO_CONDENSED_CHROMOSOME_CENTROMERIC_REGION, GO_CONDENSED_CHROMOSOME_CENTROMERIC_REGION
GO_REGULATION_OF_NUCLEAR_DIVISION, GO_REGULATION_OF_NUCLEAR_DIVISION
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME
GO_MITOTIC_CELL_CYCLE_CHECKPOINT, GO_MITOTIC_CELL_CYCLE_CHECKPOINT
GO_REGULATION_OF_GENE_SILENCING, GO_REGULATION_OF_GENE_SILENCING
GO_MEMBRANE_DOCKING, GO_MEMBRANE_DOCKING
GO_SITE_OF_DNA_DAMAGE, GO_SITE_OF_DNA_DAMAGE
GO_PML_BODY, GO_PML_BODY
GO_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_NUCLEAR_PORE, GO_NUCLEAR_PORE
GO_HETEROCHROMATIN, GO_HETEROCHROMATIN
GO_MITOTIC_SPINDLE_ORGANIZATION, GO_MITOTIC_SPINDLE_ORGANIZATION
GO_SPINDLE_ASSEMBLY, GO_SPINDLE_ASSEMBLY
GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION, GO_NUCLEAR_CHROMOSOME_TELOMERIC_REGION
GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION, GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION
GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_NEGATIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
GO_CHROMATIN_ORGANIZATION_INVOLVED_IN_REGULATION_OF_TRANSCRIPTION, GO_CHROMATIN_ORGANIZATION_INVOLVED_IN_REGULATION_OF_TRANSCRIPTION
GO_CHROMATIN_REMODELING, GO_CHROMATIN_REMODELING
GO_MITOTIC_SPINDLE, GO_MITOTIC_SPINDLE
GO_MIDBODY, GO_MIDBODY
GO_HISTONE_BINDING, GO_HISTONE_BINDING
GO_MULTI_ORGANISM_LOCALIZATION, GO_MULTI_ORGANISM_LOCALIZATION
GO_PROTEIN_LOCALIZATION_TO_CHROMATIN, GO_PROTEIN_LOCALIZATION_TO_CHROMATIN
GO_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_REGULATION_OF_ORGANELLE_ASSEMBLY
GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
GO_CHROMOSOME_SEPARATION, GO_CHROMOSOME_SEPARATION
GO_ATP_DEPENDENT_CHROMATIN_REMODELING, GO_ATP_DEPENDENT_CHROMATIN_REMODELING
GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ENDOPLASMIC_RETICULUM, GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ENDOPLASMIC_RETICULUM
GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
GO_PROTEIN_ACETYLATION, GO_PROTEIN_ACETYLATION
GO_HOST_CELLULAR_COMPONENT, GO_HOST_CELLULAR_COMPONENT
GO_SPINDLE_MICROTUBULE, GO_SPINDLE_MICROTUBULE
GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION, GO_REGULATION_OF_SISTER_CHROMATID_SEGREGATION
GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION
GO_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GO_MODIFICATION_DEPENDENT_PROTEIN_BINDING
GO_MICROTUBULE_ANCHORING, GO_MICROTUBULE_ANCHORING
GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_NUCLEUS_ORGANIZATION, GO_NUCLEUS_ORGANIZATION
GO_NUCLEAR_HETEROCHROMATIN, GO_NUCLEAR_HETEROCHROMATIN
GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS, GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS
GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS, GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS
GO_CELLULAR_PROTEIN_CONTAINING_COMPLEX_LOCALIZATION, GO_CELLULAR_PROTEIN_CONTAINING_COMPLEX_LOCALIZATION
GO_MEIOTIC_CELL_CYCLE_PROCESS, GO_MEIOTIC_CELL_CYCLE_PROCESS
GO_CHROMOSOME_LOCALIZATION, GO_CHROMOSOME_LOCALIZATION
GO_NUCLEAR_PORE_ORGANIZATION, GO_NUCLEAR_PORE_ORGANIZATION
GO_PROTEIN_METHYLATION, GO_PROTEIN_METHYLATION
GO_ANAPHASE_PROMOTING_COMPLEX, GO_ANAPHASE_PROMOTING_COMPLEX
GO_REGULATION_OF_CHROMOSOME_SEPARATION, GO_REGULATION_OF_CHROMOSOME_SEPARATION
GO_CENTROSOME_DUPLICATION, GO_CENTROSOME_DUPLICATION
GO_METAPHASE_PLATE_CONGRESSION, GO_METAPHASE_PLATE_CONGRESSION
GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON, GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON
GO_PROTEIN_N_TERMINUS_BINDING, GO_PROTEIN_N_TERMINUS_BINDING
GO_REGULATION_OF_SISTER_CHROMATID_COHESION, GO_REGULATION_OF_SISTER_CHROMATID_COHESION
GO_NEURON_PROJECTION_CYTOPLASM, GO_NEURON_PROJECTION_CYTOPLASM
GO_SPINDLE_MIDZONE_ASSEMBLY, GO_SPINDLE_MIDZONE_ASSEMBLY
GO_PEPTIDYL_LYSINE_METHYLATION, GO_PEPTIDYL_LYSINE_METHYLATION
GO_PROTEIN_LOCALIZATION_TO_MICROTUBULE_ORGANIZING_CENTER, GO_PROTEIN_LOCALIZATION_TO_MICROTUBULE_ORGANIZING_CENTER
GO_NCRNA_EXPORT_FROM_NUCLEUS, GO_NCRNA_EXPORT_FROM_NUCLEUS
GO_PORE_COMPLEX_ASSEMBLY, GO_PORE_COMPLEX_ASSEMBLY
GO_REGULATION_OF_CENTROSOME_CYCLE, GO_REGULATION_OF_CENTROSOME_CYCLE
GO_PROTEIN_LOCALIZATION_TO_KINETOCHORE, GO_PROTEIN_LOCALIZATION_TO_KINETOCHORE
GO_MITOTIC_SPINDLE_POLE, GO_MITOTIC_SPINDLE_POLE
GO_MAINTENANCE_OF_SISTER_CHROMATID_COHESION, GO_MAINTENANCE_OF_SISTER_CHROMATID_COHESION
GO_RESPONSE_TO_HYDROXYUREA, GO_RESPONSE_TO_HYDROXYUREA
GO_SEX_CHROMOSOME, GO_SEX_CHROMOSOME
GO_MITOTIC_METAPHASE_PLATE_CONGRESSION, GO_MITOTIC_METAPHASE_PLATE_CONGRESSION
GO_SISTER_CHROMATID_COHESION, GO_SISTER_CHROMATID_COHESION
GO_MITOTIC_SPINDLE_ASSEMBLY, GO_MITOTIC_SPINDLE_ASSEMBLY
GO_ORIGIN_RECOGNITION_COMPLEX, GO_ORIGIN_RECOGNITION_COMPLEX
GO_HETEROCHROMATIN_ORGANIZATION, GO_HETEROCHROMATIN_ORGANIZATION
GO_CONDENSED_NUCLEAR_CHROMOSOME_CENTROMERIC_REGION, GO_CONDENSED_NUCLEAR_CHROMOSOME_CENTROMERIC_REGION
GO_CHROMOSOME_CONDENSATION, GO_CHROMOSOME_CONDENSATION
GO_PEPTIDYL_LYSINE_DIMETHYLATION, GO_PEPTIDYL_LYSINE_DIMETHYLATION
GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
GO_MITOTIC_CYTOKINESIS, GO_MITOTIC_CYTOKINESIS
GO_REGULATION_OF_MITOTIC_SPINDLE_ASSEMBLY, GO_REGULATION_OF_MITOTIC_SPINDLE_ASSEMBLY
GO_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION, GO_REGULATION_OF_PROTEIN_AUTOPHOSPHORYLATION
GO_REGULATION_OF_SPINDLE_ORGANIZATION, GO_REGULATION_OF_SPINDLE_ORGANIZATION
GO_PEPTIDYL_LYSINE_TRIMETHYLATION, GO_PEPTIDYL_LYSINE_TRIMETHYLATION
GO_MICROTUBULE_END, GO_MICROTUBULE_END
GO_REGULATION_OF_SPINDLE_ASSEMBLY, GO_REGULATION_OF_SPINDLE_ASSEMBLY
GO_MICROTUBULE_ORGANIZING_CENTER_LOCALIZATION, GO_MICROTUBULE_ORGANIZING_CENTER_LOCALIZATION
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION
GO_MEIOSIS_II_CELL_CYCLE_PROCESS, GO_MEIOSIS_II_CELL_CYCLE_PROCESS
GO_LYSINE_N_METHYLTRANSFERASE_ACTIVITY, GO_LYSINE_N_METHYLTRANSFERASE_ACTIVITY
GO_METHYLATION_DEPENDENT_CHROMATIN_SILENCING, GO_METHYLATION_DEPENDENT_CHROMATIN_SILENCING
GO_KINETOCHORE_ASSEMBLY, GO_KINETOCHORE_ASSEMBLY
GO_A_BAND, GO_A_BAND
GO_MICROTUBULE_NUCLEATION, GO_MICROTUBULE_NUCLEATION
GO_POSITIVE_REGULATION_OF_CHROMATIN_SILENCING, GO_POSITIVE_REGULATION_OF_CHROMATIN_SILENCING
GO_POSITIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GO_POSITIVE_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION
GO_MITOTIC_SISTER_CHROMATID_COHESION, GO_MITOTIC_SISTER_CHROMATID_COHESION
GO_KINESIN_COMPLEX, GO_KINESIN_COMPLEX
GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
GO_CYTOKINESIS, GO_CYTOKINESIS
GO_PROTEIN_SERINE_THREONINE_TYROSINE_KINASE_ACTIVITY, GO_PROTEIN_SERINE_THREONINE_TYROSINE_KINASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CA

TROMERIC_REGION, GO_CHROMOSOME_CENTROMERIC_REGION