TROMERIC_REGION, GO_CHROMOSOME_CENTROMERIC_REGION

GO_KINETOCHORE_ASSEMBLY, GO_KINETOCHORE_ASSEMBLY GO_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS, GO_NUCLEOSIDE_DIPHOSPHATE_METABOLIC_PROCESS GO_LYSINE_N_METHYLTRANSFERASE_ACTIVITY, GO_LYSINE_N_METHYLTRANSFERASE_ACTIVITY GO_SPINDLE_LOCALIZATION, GO_SPINDLE_LOCALIZATION GO_NUCLEAR_PORE_OUTER_RING, GO_NUCLEAR_PORE_OUTER_RING GO_ESTABLISHMENT_OF_CENTROSOME_LOCALIZATION, GO_ESTABLISHMENT_OF_CENTROSOME_LOCALIZATION GO_ORIGIN_RECOGNITION_COMPLEX, GO_ORIGIN_RECOGNITION_COMPLEX GO_ADP_METABOLIC_PROCESS, GO_ADP_METABOLIC_PROCESS GO_CONDENSIN_COMPLEX, GO_CONDENSIN_COMPLEX GO_PYRUVATE_METABOLIC_PROCESS, GO_PYRUVATE_METABOLIC_PROCESS GO_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_PROTEOLYSIS_INVOLVED_IN_CELLULAR_PROTEIN_CATABOLIC_PROCESS GO_SEPTIN_CYTOSKELETON, GO_SEPTIN_CYTOSKELETON GO_REGULATION_OF_CYTOKINESIS, GO_REGULATION_OF_CYTOKINESIS GO_MITOTIC_SPINDLE_POLE, GO_MITOTIC_SPINDLE_POLE GO_MITOTIC_CHROMOSOME_CONDENSATION, GO_MITOTIC_CHROMOSOME_CONDENSATION GO_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC GO NUCLEOTIDE PHOSPHORYLATION, GO NUCLEOTIDE PHOSPHORYLATION GO_MEMBRANE_DISASSEMBLY, GO_MEMBRANE_DISASSEMBLY GO_PROTEIN_LOCALIZATION_TO_MICROTUBULE_ORGANIZING_CENTER, GO_PROTEIN_LOCALIZATION_TO_MICROTUBULE_ORGANIZING_CENTER GO_REGULATION_OF_SPINDLE_ORGANIZATION, GO_REGULATION_OF_SPINDLE_ORGANIZATION GO_HISTONE_METHYLTRANSFERASE_ACTIVITY, GO_HISTONE_METHYLTRANSFERASE_ACTIVITY GO_CYTOPLASMIC_DYNEIN_COMPLEX, GO_CYTOPLASMIC_DYNEIN_COMPLEX GO_MICROTUBULE_ASSOCIATED_COMPLEX, GO_MICROTUBULE_ASSOCIATED_COMPLEX

GO ANTIGEN PROCESSING AND PRESENTATION OF PEPTIDE OR POLYSACCHARIDE ANTIGEN VIA MHC CLASS II, GO ANTIGEN PROCESSING AND PRESENTATION OF PEPTIDE OR POLYSACCHARIDE ANTIGEN.

GO_PHOSPHATASE_INHIBITOR_ACTIVITY, GO_PHOSPHATASE_INHIBITOR_ACTIVITY

GO_PROMOTER_SPECIFIC_CHROMATIN_BINDING, GO_PROMOTER_SPECIFIC_CHROMATIN_BINDING

GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON, GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON

GO_REGULATION_OF_MITOTIC_SPINDLE_ASSEMBLY, GO_REGULATION_OF_MITOTIC_SPINDLE_ASSEMBLY

GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS, GO_REGULATION_OF_CARBOHYDRATE_CATABOLIC_PROCESS

GO_CONDENSED_CHROMOSOME_OUTER_KINETOCHORE, GO_CONDENSED_CHROMOSOME_OUTER_KINETOCHORE

GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT

GO_CARBOHYDRATE_DERIVATIVE_CATABOLIC_PROCESS, GO_CARBOHYDRATE_DERIVATIVE_CATABOLIC_PROCESS

GO_REGULATION_OF_SPINDLE_CHECKPOINT, GO_REGULATION_OF_SPINDLE_CHECKPOINT

GO_POSITIVE_REGULATION_OF_CYTOKINESIS, GO_POSITIVE_REGULATION_OF_CYTOKINESIS

GO_REGULATION_OF_CENTROSOME_CYCLE, GO_REGULATION_OF_CENTROSOME_CYCLE

GO_FEMALE_MEIOTIC_NUCLEAR_DIVISION, GO_FEMALE_MEIOTIC_NUCLEAR_DIVISION

GO_REGULATION_OF_HISTONE_METHYLATION, GO_REGULATION_OF_HISTONE_METHYLATION

GO_MAINTENANCE_OF_LOCATION_IN_CELL, GO_MAINTENANCE_OF_LOCATION_IN_CELL

GO_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX, GO_PROTEIN_PHOSPHATASE_TYPE_2A_COMPLEX

GO_ANAPHASE_PROMOTING_COMPLEX, GO_ANAPHASE_PROMOTING_COMPLEX

GO_CHROMOSOME_CONDENSATION, GO_CHROMOSOME_CONDENSATION

GO_BASEMENT_MEMBRANE_ASSEMBLY, GO_BASEMENT_MEMBRANE_ASSEMBLY

GO_MALE_GERM_CELL_NUCLEUS, GO_MALE_GERM_CELL_NUCLEUS

GO_SEH1_ASSOCIATED_COMPLEX, GO_SEH1_ASSOCIATED_COMPLEX

GO_EXIT_FROM_MITOSIS, GO_EXIT_FROM_MITOSIS