

TROMERIC_REGION, GO_CHROMOSOME_CENTROMERIC_REGION

- GO_SPINDLE_ORGANIZATION, GO_SPINDLE_ORGANIZATION
- GO_KINETOCHORE, GO_KINETOCHORE
- GO_HISTONE_BINDING, GO_HISTONE_BINDING
- GO_SPINDLE_POLE, GO_SPINDLE_POLE
- GO_MIDBODY, GO_MIDBODY
- GO_CONDENSED_CHROMOSOME_CENTROMERIC_REGION, GO_CONDENSED_CHROMOSOME_CENTROMERIC_REGION
- GO_SPINDLE_ASSEMBLY, GO_SPINDLE_ASSEMBLY
- GO_MODIFICATION_DEPENDENT_PROTEIN_BINDING, GO_MODIFICATION_DEPENDENT_PROTEIN_BINDING
- GO_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS, GO_MICROTUBULE_CYTOSKELETON_ORGANIZATION_INVOLVED_IN_MITOSIS
- GO_PROTEIN_METHYLATION, GO_PROTEIN_METHYLATION
- GO_MITOTIC_SPINDLE, GO_MITOTIC_SPINDLE
- GO_MITOTIC_SPINDLE_ORGANIZATION, GO_MITOTIC_SPINDLE_ORGANIZATION
- GO_PEPTIDYL_LYSINE_METHYLATION, GO_PEPTIDYL_LYSINE_METHYLATION
- GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
- GO_METHYLATED_HISTONE_BINDING, GO_METHYLATED_HISTONE_BINDING
- GO_SPINDLE_MICROTUBULE, GO_SPINDLE_MICROTUBULE
- GO_ANAPHASE_PROMOTING_COMPLEX_DEPENDENT_CATABOLIC_PROCESS, GO_ANAPHASE_PROMOTING_COMPLEX_DEPENDENT_CATABOLIC_PROCESS
- GO_CILIARY_BASAL_BODY_PLASMA_MEMBRANE_DOCKING, GO_CILIARY_BASAL_BODY_PLASMA_MEMBRANE_DOCKING
- GO_MEMBRANE_DOCKING, GO_MEMBRANE_DOCKING
- GO_HISTONE_METHYLATION, GO_HISTONE_METHYLATION
- GO_NUCLEAR_PORE, GO_NUCLEAR_PORE
- GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ENDOPLASMIC_RETICULUM, GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ENDOPLASMIC_RETICULUM
- GO_REGULATION_OF_CHROMOSOME_SEPARATION, GO_REGULATION_OF_CHROMOSOME_SEPARATION
- GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
- GO_MULTI_ORGANISM_LOCALIZATION, GO_MULTI_ORGANISM_LOCALIZATION
- GO_PEPTIDYL_LYSINE_TRIMETHYLATION, GO_PEPTIDYL_LYSINE_TRIMETHYLATION
- GO_NCRNA_EXPORT_FROM_NUCLEUS, GO_NCRNA_EXPORT_FROM_NUCLEUS
- GO_PHOSPHATASE_COMPLEX, GO_PHOSPHATASE_COMPLEX
- GO_CYTOKINESIS, GO_CYTOKINESIS
- GO_NUCLEUS_ORGANIZATION, GO_NUCLEUS_ORGANIZATION
- GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION, GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION
- GO_HOST_CELLULAR_COMPONENT, GO_HOST_CELLULAR_COMPONENT
- GO_METAPHASE_PLATE_CONGRESSION, GO_METAPHASE_PLATE_CONGRESSION
- GO_PROTEIN_SUMOYLATION, GO_PROTEIN_SUMOYLATION
- GO_CHROMOSOME_LOCALIZATION, GO_CHROMOSOME_LOCALIZATION
- GO_MITOTIC_SPINDLE_ASSEMBLY, GO_MITOTIC_SPINDLE_ASSEMBLY
- GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GO_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
- GO_MICROTUBULE_NUCLEATION, GO_MICROTUBULE_NUCLEATION
- GO_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_REGULATION_OF_ORGANELLE_ASSEMBLY
- GO_MITOTIC_CHROMOSOME_CONDENSATION, GO_MITOTIC_CHROMOSOME_CONDENSATION
- GO_HISTONE_METHYLTRANSFERASE_ACTIVITY, GO_HISTONE_METHYLTRANSFERASE_ACTIVITY
- GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS, GO_REGULATION_OF_PROTEIN_EXPORT_FROM_NUCLEUS
- GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON, GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON
- GO_REGULATION_OF_CHROMATIN_BINDING, GO_REGULATION_OF_CHROMATIN_BINDING
- GO_PERICENTRIC_HETEROCHROMATIN, GO_PERICENTRIC_HETEROCHROMATIN
- GO_LYSINE_N_METHYLTRANSFERASE_ACTIVITY, GO_LYSINE_N_METHYLTRANSFERASE_ACTIVITY
- GO_REGULATION_OF_CYTOKINESIS, GO_REGULATION_OF_CYTOKINESIS
- GO_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC, GO_POSITIVE_REGULATION_OF_GENE_EXPRESSION_EPIGENETIC
- GO_HISTONE_H4_K5_ACETYLTATION, GO_HISTONE_H4_K5_ACETYLTATION
- GO_CARBOHYDRATE_CATABOLIC_PROCESS, GO_CARBOHYDRATE_CATABOLIC_PROCESS
- GO_ESTABLISHMENT_OF_CELL_POLARITY, GO_ESTABLISHMENT_OF_CELL_POLARITY
- GO_MITOTIC_METAPHASE_PLATE_CONGRESSION, GO_MITOTIC_METAPHASE_PLATE_CONGRESSION
- GO_MICROTUBULE_PLUS_END, GO_MICROTUBULE_PLUS_END
- GO_HISTONE_LYSINE_N_METHYLTRANSFERASE_ACTIVITY, GO_HISTONE_LYSINE_N_METHYLTRANSFERASE_ACTIVITY
- GO_MICROTUBULE_ASSOCIATED_COMPLEX, GO_MICROTUBULE_ASSOCIATED_COMPLEX
- GO_REGULATION_OF_SPINDLE_ASSEMBLY, GO_REGULATION_OF_SPINDLE_ASSEMBLY
- GO_PROTEIN_PHOSPHATASE_2A_BINDING, GO_PROTEIN_PHOSPHATASE_2A_BINDING
- GO_CYTOSKELETON_DEPENDENT_CYTOKINESIS, GO_CYTOSKELETON_DEPENDENT_CYTOKINESIS
- GO_KINETOCHORE_ORGANIZATION, GO_KINETOCHORE_ORGANIZATION
- GO_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE
- GO_MICROTUBULE_ORGANIZING_CENTER_LOCALIZATION, GO_MICROTUBULE_ORGANIZING_CENTER_LOCALIZATION
- GO_POSITIVE_REGULATION_OF_TORC1_SIGNALING, GO_POSITIVE_REGULATION_OF_TORC1_SIGNALING
- GO_POSITIVE_REGULATION_OF_CHROMATIN_BINDING, GO_POSITIVE_REGULATION_OF_CHROMATIN_BINDING
- GO_MICROTUBULE_PLUS_END_BINDING, GO_MICROTUBULE_PLUS_END_BINDING
- GO_PROTEIN_LOCALIZATION_TO_MICROTUBULE_ORGANIZING_CENTER, GO_PROTEIN_LOCALIZATION_TO_MICROTUBULE_ORGANIZING_CENTER