

	GO_TELOMERE_ORGANIZATION, GO_TELOMERE_ORGANIZATION
	GO_CHROMOSOME_CENTROMERIC_REGION, GO_CHROMOSOME_CENTROMERIC_REGION
	GO_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_MITOTIC_SISTER_CHROMATID_SEGREGATION
	GO_DNA_DEPENDENT_DNA_REPLICATION, GO_DNA_DEPENDENT_DNA_REPLICATION
	GO_DNA_DEPENDENT_ATPASE_ACTIVITY, GO_DNA_DEPENDENT_ATPASE_ACTIVITY
	GO_DNA_PACKAGING_COMPLEX, GO_DNA_PACKAGING_COMPLEX
	GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
	GO_SPINDLE_ORGANIZATION, GO_SPINDLE_ORGANIZATION
	GO_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_REGULATION_OF_CHROMOSOME_SEGREGATION
	GO_MIDBODY, GO_MIDBODY
	GO_SPINDLE_POLE, GO_SPINDLE_POLE
	GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_ORGANIZATION
	GO_RECOMBINATIONAL_REPAIR, GO_RECOMBINATIONAL_REPAIR
	GO_ANAPHASE_PROMOTING_COMPLEX_DEPENDENT_CATABOLIC_PROCESS, GO_ANAPHASE_PROMOTING_COMPLEX_DEPENDENT_CATABOLIC_PROCESS
	GO_MITOTIC_CELL_CYCLE_CHECKPOINT, GO_MITOTIC_CELL_CYCLE_CHECKPOINT
	GO_SPINDLE_ASSEMBLY, GO_SPINDLE_ASSEMBLY
	GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
	GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
	GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME
	GO_SITE_OF_DNA_DAMAGE, GO_SITE_OF_DNA_DAMAGE
	GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
	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_CENTROMERE_COMPLEX_ASSEMBLY, GO_CENTROMERE_COMPLEX_ASSEMBLY
	GO_CILIARY_BASAL_BODY_PLASMA_MEMBRANE_DOCKING, GO_CILIARY_BASAL_BODY_PLASMA_MEMBRANE_DOCKING
	GO_CELL_CYCLE_DNA_REPLICATION, GO_CELL_CYCLE_DNA_REPLICATION
	GO_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX, GO_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX
	GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION
	GO_SITE_OF_DOUBLE_STRAND_BREAK, GO_SITE_OF_DOUBLE_STRAND_BREAK
	GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
	GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS, GO_RNA_DEPENDENT_DNA_BIOSYNTHETIC_PROCESS
	GO_SISTER_CHROMATID_COHESION, GO_SISTER_CHROMATID_COHESION
	GO_REGULATION_OF_DNA_REPAIR, GO_REGULATION_OF_DNA_REPAIR
	GO_HETEROCHROMATIN, GO_HETEROCHROMATIN
	GO_CHROMOSOME_SEPARATION, GO_CHROMOSOME_SEPARATION
	GO_REGULATION_OF_DNA_REPLICATION, GO_REGULATION_OF_DNA_REPLICATION
	GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION, GO_MICROTUBULE_ORGANIZING_CENTER_ORGANIZATION
	GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN
	GO_REGULATION_OF_CHROMATIN_ORGANIZATION, GO_REGULATION_OF_CHROMATIN_ORGANIZATION
	GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_POSITIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
	GO_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_REGULATION_OF_ORGANELLE_ASSEMBLY
	GO_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION, GO_REGULATION_OF_DNA_DEPENDENT_DNA_REPLICATION
	GO_DNA_BIOSYNTHETIC_PROCESS, GO_DNA_BIOSYNTHETIC_PROCESS
	GO_REGULATION_OF_CHROMOSOME_SEPARATION, GO_REGULATION_OF_CHROMOSOME_SEPARATION
	GO_MITOTIC_SPINDLE, GO_MITOTIC_SPINDLE
	GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY, GO_DNA_DEPENDENT_DNA_REPLICATION_MAINTENANCE_OF_FIDELITY
	GO_CYTOKINESIS, GO_CYTOKINESIS
	GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT, GO_MITOTIC_DNA_INTEGRITY_CHECKPOINT
	GO_REGULATION_OF_SPINDLE_ASSEMBLY, GO_REGULATION_OF_SPINDLE_ASSEMBLY
	GO_TELOMERE_CAPPING, GO_TELOMERE_CAPPING
	GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_SPINDLE_MICROTUBULES_TO_KINETOCHORE
	GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS, GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS
	GO_CYTOSKELETON_DEPENDENT_CYTOKINESIS, GO_CYTOSKELETON_DEPENDENT_CYTOKINESIS
	GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
	GO_SPINDLE_MICROTUBULE, GO_SPINDLE_MICROTUBULE
	GO_REGULATION_OF_CENTROSOME_CYCLE, GO_REGULATION_OF_CENTROSOME_CYCLE
	GO_MITOTIC_CYTOKINESIS, GO_MITOTIC_CYTOKINESIS
	GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GO_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
	GO_REPLICATION_FORK_PROCESSING, GO_REPLICATION_FORK_PROCESSING
	GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_NEGATIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
	GO_LATE_ENDOSOME_MEMBRANE, GO_LATE_ENDOSOME_MEMBRANE
	GO_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_DNA_METABOLIC_PROCESS
	GO_REGULATION_OF_NUCLEAR_DIVISION, GO_REGULATION_OF_NUCLEAR_DIVISION
	GO_CYTOKINETIC_PROCESS, GO_CYTOKINETIC_PROCESS
	GO_NUCLEAR_HETEROCHROMATIN, GO_NUCLEAR_HETEROCHROMATIN
	GO_ENDOSOME_ORGANIZATION, GO_ENDOSOME_ORGANIZATION
	GO_METAPHASE_PLATE_CONGRESSION, GO_METAPHASE_PLATE_CONGRESSION
	GO_FLEMMING_BODY, GO_FLEMMING_BODY
	GO_HISTONE_DEACETYLATION, GO_HISTONE_DEACETYLATION
	GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE, GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE
	GO_MITOTIC_METAPHASE_PLATE_CONGRESSION, GO_MITOTIC_METAPHASE_PLATE_CONGRESSION
	GO_MICROTUBULE_BUNDLE_FORMATION, GO_MICROTUBULE_BUNDLE_FORMATION
	GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_DNA_BIOSYNTHETIC_PROCESS
	GO_REGULATION_OF_CENTROSOME_DUPLICATION, GO_REGULATION_OF_CENTROSOME_DUPLICATION
	GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING, GO_NEGATIVE_REGULATION_OF_TELOMERE_MAINTENANCE_VIA_TELOMERE_LENGTHENING
	GO_VACUOLAR_TRANSPORT, GO_VACUOLAR_TRANSPORT
	GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_TELOMERIC_REGION
	GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME, GO_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_CHROMOSOME
	GO_PROTEIN_C_TERMINUS_BINDING, GO_PROTEIN_C_TERMINUS_BINDING
	GO_REGULATION_OF_SISTER_CHROMATID_COHESION, GO_REGULATION_OF_SISTER_CHROMATID_COHESION
	GO_POSITIVE_REGULATION_OF_DNA_REPAIR, GO_POSITIVE_REGULATION_OF_DNA_REPAIR
	GO_NEGATIVE_REGULATION_OF_DNA_REPLICATION, GO_NEGATIVE_REGULATION_OF_DNA_REPLICATION
	GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX, GO_NUCLEAR_UBIQUITIN_LIGASE_COMPLEX
	GO_REGULATION_OF_MITOTIC_SPINDLE_ASSEMBLY, GO_REGULATION_OF_MITOTIC_SPINDLE_ASSEMBLY
	GO_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR, GO_POSITIVE_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR
	GO_MULTIVESICULAR_BODY_ORGANIZATION, GO_MULTIVESICULAR_BODY_ORGANIZATION
	GO_MITOTIC_SPINDLE_ASSEMBLY, GO_MITOTIC_SPINDLE_ASSEMBLY
	GO_MITOTIC_SISTER_CHROMATID_COHESION, GO_MITOTIC_SISTER_CHROMATID_COHESION
	GO_NEGATIVE_REGULATION_OF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE, GO_NEGATIVE_REGULATION_OF_METAPHASE_ANAPHASE_TRANSITION_OF_CELL_CYCLE
	GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION, GO_PROTEIN_LOCALIZATION_TO_CHROMOSOME_CENTROMERIC_REGION
	GO_PROTEIN_LOCALIZATION_TO_CHROMATIN, GO_PROTEIN_LOCALIZATION_TO_CHROMATIN
	GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_TELOMERASE_ACTIVITY
	GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ENDOPLASMIC_RETICULUM, GO_RETROGRADE_VESICLE_MEDIATED_TRANSPORT_GOLGI_TO_ENDOPLASMIC_RETICULUM
	GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_POSITIVE_REGULATION_OF_MITOTIC_SISTER_CHROMATID_SEGREGATION
	GO_FEMALE_MEIOTIC_NUCLEAR_DIVISION, GO_FEMALE_MEIOTIC_NUCLEAR_DIVISION
	GO_NUCLEAR_HORMONE_RECEPTOR_BINDING, GO_NUCLEAR_HORMONE_RECEPTOR_BINDING
	GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DNA_DAMAGE_STIMULUS
	GO_MITOTIC_SPINDLE_POLE, GO_MITOTIC_SPINDLE_POLE
	GO_NEGATIVE_REGULATION_OF_NUCLEAR_DIVISION, GO_NEGATIVE_REGULATION_OF_NUCLEAR_DIVISION
	GO_G2_DNA_DAMAGE_CHECKPOINT, GO_G2_DNA_DAMAGE_CHECKPOINT
	GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY, GO_SINGLE_STRANDED_DNA_HELICASE_ACTIVITY
	GO_NEGATIVE_REGULATION_OF_ORGANELLE_ASSEMBLY, GO_NEGATIVE_REGULATION_OF_ORGANELLE_ASSEMBLY
	GO_REGULATION_OF_HISTONE_DEACETYLATION, GO_REGULATION_OF_HISTONE_DEACETYLATION
	GO_DEOXYRIBONUCLEASE_ACTIVITY, GO_DEOXYRIBONUCLEASE_ACTIVITY
	GO_CENTROSOME_DUPLICATION, GO_CENTROSOME_DUPLICATION
	GO_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE, GO_ATTACHMENT_OF_MITOTIC_SPINDLE_MICROTUBULES_TO_KINETOCHORE
	GO_REGULATION_OF_TELOMERE_CAPPING, GO_REGULATION_OF_TELOMERE_CAPPING
	GO_ANAPHASE_PROMOTING_COMPLEX, GO_ANAPHASE_PROMOTING_COMPLEX
	GO_CYTOPLASMIC_DYNEIN_COMPLEX, GO_CYTOPLASMIC_DYNEIN_COMPLEX
	GO_DNA_SECONDARY_STRUCTURE_BINDING, GO_DNA_SECONDARY_STRUCTURE_BINDING
	GO_SPINDLE_LOCALIZATION, GO_SPINDLE_LOCALIZATION
	GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON, GO_PROTEIN_LOCALIZATION_TO_CYTOSKELETON
	GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY, GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY
	GO_KINETOCHORE_ORGANIZATION, GO_KINETOCHORE_ORGANIZATION
	GO_PROTEIN_LOCALIZATION_TO_KINETOCHORE, GO_PROTEIN_LOCALIZATION_TO_KINETOCHORE
	GO_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I, GO_NUCLEOLAR_LARGE_RRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_I
	GO_FOUR_WAY_JUNCTION_DNA_BINDING, GO_FOUR_WAY_JUNCTION_DNA_BINDING
	GO_MICROTUBULE_ASSOCIATED_COMPLEX, GO_MICROTUBULE_ASSOCIATED_COMPLEX
	GO_MALE_MEIOTIC_NUCLEAR_DIVISION, GO_MALE_MEIOTIC_NUCLEAR_DIVISION
	GO_FLAP_ENDONUCLEASE_ACTIVITY, GO_FLAP_ENDONUCLEASE_ACTIVITY
	GO_NEGATIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
	GO_SPINDLE_MIDZONE, GO_SPINDLE_MIDZONE