
GRITY_CHECKPOINT, GO_DNA_INTEGRITY_CHECKPOINT
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GO_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
GO_POSITIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION, GO_POSITIVE_REGULATION_OF_ESTABLISHMENT_OF_PROTEIN_LOCALIZATION_TO_MITOCHONDRION
GO_SERINE_THREONINE_PROTEIN_KINASE_COMPLEX, GO_SERINE_THREONINE_PROTEIN_KINASE_COMPLEX
GO_PROTEIN_KINASE_COMPLEX, GO_PROTEIN_KINASE_COMPLEX
GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE, GO_SIGNAL_TRANSDUCTION_IN_RESPONSE_TO_DNA_DAMAGE
GO_RESPONSE_TO_UV, GO_RESPONSE_TO_UV
GO_U2_TYPE_CATALYTIC_STEP_2_SPLICEOSOME, GO_U2_TYPE_CATALYTIC_STEP_2_SPLICEOSOME
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_BY_P53_CLASS_MEDIATOR
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE_BY_P53_CLASS_MEDIATOR, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE_BY_P53
GO_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_REGULATOR_ACTIVITY, GO_CYCLIN_DEPENDENT_PROTEIN_SERINE_THREONINE_KINASE_REGULATOR_ACTIVITY
GO_POSITIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL, GO_POSITIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION
GO_NEGATIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_P
GO_POSITIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_CYCLIN_DEPENDENT_PROTEIN_KINASE_ACTIVITY
GO_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX, GO_CYCLIN_DEPENDENT_PROTEIN_KINASE_HOLOENZYME_COMPLEX
GO_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING, GO_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_POLY_A_TAIL_SHORTENING
GO_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX, GO_CUL4_RING_E3_UBIQUITIN_LIGASE_COMPLEX
GO_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENYLATION_DEPENDENT_DECAY, GO_REGULATION_OF_NUCLEAR_TRANSCRIBED_MRNA_CATABOLIC_PROCESS_DEADENY
GO_PROTEIN_METHYLTRANSFERASE_ACTIVITY, GO_PROTEIN_METHYLTRANSFERASE_ACTIVITY
GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING, GO_REGULATION_OF_DOUBLE_STRAND_BREAK_REPAIR_VIA_NONHOMOLOGOUS_END_JOINING
GO_REGULATION_OF_TYPE_I_INTERFERON_MEDIATED_SIGNALING_PATHWAY, GO_REGULATION_OF_TYPE_I_INTERFERON_MEDIATED_SIGNALING_PATHWAY
GO_CELLULAR_RESPONSE_TO_UV, GO_CELLULAR_RESPONSE_TO_UV
GO_ORGANELLE_INHERITANCE, GO_ORGANELLE_INHERITANCE
GO_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE, GO_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE
GO_SIGNAL_TRANSDUCTION_INVOLVED_IN_CELL_CYCLE_CHECKPOINT, GO_SIGNAL_TRANSDUCTION_INVOLVED_IN_CELL_CYCLE_CHECKPOINT
GO_ENTRAINMENT_OF_CIRCADIAN_CLOCK, GO_ENTRAINMENT_OF_CIRCADIAN_CLOCK
GO_POSITIVE_REGULATION_OF_VIRAL_GENOME_REPLICATION, GO_POSITIVE_REGULATION_OF_VIRAL_GENOME_REPLICATION
GO_HISTONE_METHYLTRANSFERASE_ACTIVITY, GO_HISTONE_METHYLTRANSFERASE_ACTIVITY
GO_PHOSPHATASE_BINDING, GO_PHOSPHATASE_BINDING
GO_BLASTOCYST_DEVELOPMENT, GO_BLASTOCYST_DEVELOPMENT
GO_SCF_UBIQUITIN_LIGASE_COMPLEX, GO_SCF_UBIQUITIN_LIGASE_COMPLEX
GO_MITOCHONDRIAL_DNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_DNA_METABOLIC_PROCESS
GO_KINASE_INHIBITOR_ACTIVITY, GO_KINASE_INHIBITOR_ACTIVITY
GO_PHOTOPERIODISM, GO_PHOTOPERIODISM
GO_ATRIOVENTRICULAR_VALVE_DEVELOPMENT, GO_ATRIOVENTRICULAR_VALVE_DEVELOPMENT
GO_CORTICOSTEROID_RECEPTOR_SIGNALING_PATHWAY, GO_CORTICOSTEROID_RECEPTOR_SIGNALING_PATHWAY