
OSTTRANSCRIPTIONAL_GENE_SILENCING, GO_RNA_BINDING_INVOLVED_IN_POSTTRANSCRIPTIONAL_GENE_SILENCING
GO_MIRNA_MEDIATED_INHIBITION_OF_TRANSLATION, GO_MIRNA_MEDIATED_INHIBITION_OF_TRANSLATION
GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING, GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING
GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT
GO_NEUROINFLAMMATORY_RESPONSE, GO_NEUROINFLAMMATORY_RESPONSE
GO_REGULATION_OF_MUSCLE_CONTRACTION, GO_REGULATION_OF_MUSCLE_CONTRACTION
GO_RNA_POLYMERASE_BINDING, GO_RNA_POLYMERASE_BINDING
GO_CARDIOCYTE_DIFFERENTIATION, GO_CARDIOCYTE_DIFFERENTIATION
GO_RNA_POLYMERASE_CORE_ENZYME_BINDING, GO_RNA_POLYMERASE_CORE_ENZYME_BINDING
GO_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GO_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION
GO_FORMATION_OF_PRIMARY_GERM_LAYER, GO_FORMATION_OF_PRIMARY_GERM_LAYER
GO_ACTOMYOSIN_STRUCTURE_ORGANIZATION, GO_ACTOMYOSIN_STRUCTURE_ORGANIZATION
GO_REGULATION_OF_POTASSIUM_ION_TRANSPORT, GO_REGULATION_OF_POTASSIUM_ION_TRANSPORT
GO_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_CHOLESTEROL_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT, GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT
GO_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GO_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION
GO_STRIATED_MUSCLE_CONTRACTION, GO_STRIATED_MUSCLE_CONTRACTION
GO_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_REGULATION_OF_SPROUTING_ANGIOGENESIS
GO_ARTERY_MORPHOGENESIS, GO_ARTERY_MORPHOGENESIS
GO_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION, GO_POSITIVE_REGULATION_OF_SMOOTH_MUSCLE_CELL_PROLIFERATION
GO_SPROUTING_ANGIOGENESIS, GO_SPROUTING_ANGIOGENESIS
GO_CARDIAC_CELL_DEVELOPMENT, GO_CARDIAC_CELL_DEVELOPMENT
GO_CARDIAC_MUSCLE_CONTRACTION, GO_CARDIAC_MUSCLE_CONTRACTION
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GO_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOPMENT, GO_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOP
GO_GASTRULATION, GO_GASTRULATION
GO_ORGAN_GROWTH, GO_ORGAN_GROWTH
GO_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION, GO_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION
GO_REGULATION_OF_CELL_FATE_COMMITMENT, GO_REGULATION_OF_CELL_FATE_COMMITMENT
GO_NEGATIVE_REGULATION_OF_CHEMOTAXIS, GO_NEGATIVE_REGULATION_OF_CHEMOTAXIS
GO_NEGATIVE_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GO_NEGATIVE_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE
GO_REGULATION_OF_CELL_CHEMOTAXIS_TO_FIBROBLAST_GROWTH_FACTOR, GO_REGULATION_OF_CELL_CHEMOTAXIS_TO_FIBROBLAST_GROWTH_FACTOR
GO_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GO_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
GO_NEGATIVE_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION
GO_REGULATION_OF_MESODERM_FORMATION, GO_REGULATION_OF_MESODERM_FORMATION
GO_INTERLEUKIN_1_BIOSYNTHETIC_PROCESS, GO_INTERLEUKIN_1_BIOSYNTHETIC_PROCESS
GO_INTERLEUKIN_1_BETA_BIOSYNTHETIC_PROCESS, GO_INTERLEUKIN_1_BETA_BIOSYNTHETIC_PROCESS
GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION
GO_REGULATION_OF_CARDIAC_MUSCLE_CELL_DIFFERENTIATION, GO_REGULATION_OF_CARDIAC_MUSCLE_CELL_DIFFERENTIATION
GO_TISSUE_REGENERATION, GO_TISSUE_REGENERATION
GO_MESODERMAL_CELL_DIFFERENTIATION, GO_MESODERMAL_CELL_DIFFERENTIATION
GO_REGULATION_OF_NEUROINFLAMMATORY_RESPONSE, GO_REGULATION_OF_NEUROINFLAMMATORY_RESPONSE
GO_NEGATIVE_REGULATION_OF_DEVELOPMENTAL_GROWTH, GO_NEGATIVE_REGULATION_OF_DEVELOPMENTAL_GROWTH
GO_MESODERM_MORPHOGENESIS, GO_MESODERM_MORPHOGENESIS
GO_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION, GO_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION
GO_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS, GO_REGULATION_OF_COLLAGEN_METABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_MITOTIC_CELL_CYCLE, GO_POSITIVE_REGULATION_OF_MITOTIC_CELL_CYCLE
GO_CARDIAC_MUSCLE_CELL_DIFFERENTIATION, GO_CARDIAC_MUSCLE_CELL_DIFFERENTIATION
GO_REGULATION_OF_VASCULAR_SMOOTH_MUSCLE_CELL_PROLIFERATION, GO_REGULATION_OF_VASCULAR_SMOOTH_MUSCLE_CELL_PROLIFERATION
GO_MUSCLE_CELL_DEVELOPMENT, GO_MUSCLE_CELL_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GO_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE
GO_NEGATIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_NEGATIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS
GO_NEGATIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS, GO_NEGATIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS
GO_REGULATION_OF_CELLULAR_KETONE_METABOLIC_PROCESS, GO_REGULATION_OF_CELLULAR_KETONE_METABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_CELL_FATE_COMMITMENT, GO_POSITIVE_REGULATION_OF_CELL_FATE_COMMITMENT
GO_NEGATIVE_REGULATION_OF_STEM_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_STEM_CELL_PROLIFERATION
GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_NEGATIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
GO_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT, GO_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT
GO_MEMBRANE_REPOLARIZATION_DURING_ACTION_POTENTIAL, GO_MEMBRANE_REPOLARIZATION_DURING_ACTION_POTENTIAL
GO_MESODERM_DEVELOPMENT, GO_MESODERM_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_MYOBLAST_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_MYOBLAST_PROLIFERATION
GO_COLLAGEN_BIOSYNTHETIC_PROCESS, GO_COLLAGEN_BIOSYNTHETIC_PROCESS
GO_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY, GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORTER_ACTIVITY
GO_NEGATIVE_REGULATION_OF_OSSIFICATION, GO_NEGATIVE_REGULATION_OF_OSSIFICATION
GO_REGULATION_OF_RESPONSE_TO_WOUNDING, GO_REGULATION_OF_RESPONSE_TO_WOUNDING
GO_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_PROTEIN_CONJUGATION_OR_REMOVAL, GO_NEGATIVE_REGULATION_OF_PROTEIN_MODIFICATION_BY_SMALL_P
GO_NEGATIVE_REGULATION_OF_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_FIBROBLAST_GROWTH_FACTOR_RECEPTOR_SIGNA
GO_ADRENERGIC_RECEPTOR_SIGNALING_PATHWAY, GO_ADRENERGIC_RECEPTOR_SIGNALING_PATHWAY
GO_POSITIVE_REGULATION_OF_PROTEIN_KINASE_B_SIGNALING, GO_POSITIVE_REGULATION_OF_PROTEIN_KINASE_B_SIGNALING
GO_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GO_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
GO_PHYSIOLOGICAL_CARDIAC_MUSCLE_HYPERTROPHY, GO_PHYSIOLOGICAL_CARDIAC_MUSCLE_HYPERTROPHY
GO_PHENOTYPIC_SWITCHING, GO_PHENOTYPIC_SWITCHING
GO_POSITIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_CELLULAR_AMIDE_METABOLIC_PROCESS
GO_POSITIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_POSITIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS
GO_CONTRACTILE_ACTIN_FILAMENT_BUNDLE_ASSEMBLY, GO_CONTRACTILE_ACTIN_FILAMENT_BUNDLE_ASSEMBLY
GO_REGULATION_OF_MESODERMAL_CELL_DIFFERENTIATION, GO_REGULATION_OF_MESODERMAL_CELL_DIFFERENTIATION
GO_ARTERY_DEVELOPMENT, GO_ARTERY_DEVELOPMENT
GO_NEGATIVE_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_MUSCLE_CELL_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_CELL_GROWTH_INVOLVED_IN_CARDIAC_M