DSTTRANSCRIPTIONAL_GENE_SILENCING, GO_RNA_BINDING_INVOLVED_IN_POSTTRANSCRIPTIONAL_GENE_SILENCING

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GO POSITIVE REGULATION OF CELL CYCLE PHASE TRANSITION, GO POSITIVE REGULATION OF CELL CYCLE PHASE TRANSITION
GO_REGULATION_OF_WOUND_HEALING, GO_REGULATION_OF_WOUND_HEALING
GO_POSITIVE_REGULATION_OF_MITOTIC_CELL_CYCLE, GO_POSITIVE_REGULATION_OF_MITOTIC_CELL_CYCLE
GO_INTERLEUKIN_1_PRODUCTION, GO_INTERLEUKIN_1_PRODUCTION
GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING, GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING
GO_NEUROTRANSMITTER_METABOLIC_PROCESS, GO_NEUROTRANSMITTER_METABOLIC_PROCESS
GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT
GO_RNA_POLYMERASE_BINDING, GO_RNA_POLYMERASE_BINDING
GO_NEGATIVE_REGULATION_OF_OXIDOREDUCTASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_OXIDOREDUCTASE_ACTIVITY
GO_COLLAGEN_METABOLIC_PROCESS, GO_COLLAGEN_METABOLIC_PROCESS
GO_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GO_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
GO_ADENYLATE_CYCLASE_INHIBITING_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GO_ADENYLATE_CYCLASE_INHIBITING_G_PROTEIN_COUPLED_RECEPTOR
GO_NEGATIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSCLE_CELL_MIGRATION, GO_NEGATIVE_REGULATION_OF_VASCULAR_ASSOCIATED_SMOOTH_MUSC
GO_INTERLEUKIN_1_BETA_PRODUCTION, GO_INTERLEUKIN_1_BETA_PRODUCTION
GO_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT, GO_REGULATION_OF_CONNECTIVE_TISSUE_REPLACEMENT
GO_NEUROTRANSMITTER_BIOSYNTHETIC_PROCESS, GO_NEUROTRANSMITTER_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_REGULATION_OF_SPROUTING_ANGIOGENESIS
GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION, GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_8_PRODUCTION
GO NEGATIVE REGULATION OF INFLAMMATORY RESPONSE, GO NEGATIVE REGULATION OF INFLAMMATORY RESPONSE
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DRUG, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_DRUG
GO_NEUROINFLAMMATORY_RESPONSE, GO_NEUROINFLAMMATORY_RESPONSE
GO_TISSUE_REMODELING, GO_TISSUE_REMODELING
GO_NIK_NF_KAPPAB_SIGNALING, GO_NIK_NF_KAPPAB_SIGNALING
GO_INTERLEUKIN_6_PRODUCTION, GO_INTERLEUKIN_6_PRODUCTION
GO_POSITIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_POSITIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS
GO_POSITIVE_REGULATION_OF_TISSUE_REMODELING, GO_POSITIVE_REGULATION_OF_TISSUE_REMODELING
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_WOUNDING
GO_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GO_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE
GO_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS, GO_REGULATION_OF_ENDOTHELIAL_CELL_CHEMOTAXIS
GO_RNA_POLYMERASE_CORE_ENZYME_BINDING, GO_RNA_POLYMERASE_CORE_ENZYME_BINDING
GO_NEGATIVE_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE, GO_NEGATIVE_REGULATION_OF_LIPOPROTEIN_PARTICLE_CLEARANCE
GO_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS, GO_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS
GO_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GO_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE
GO_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE, GO_NEGATIVE_REGULATION_OF_LOW_DENSITY_LIPOPROTEIN_PARTICLE_CLEARANCE
GO_SPROUTING_ANGIOGENESIS, GO_SPROUTING_ANGIOGENESIS
GO_POSITIVE_REGULATION_OF_WOUND_HEALING, GO_POSITIVE_REGULATION_OF_WOUND_HEALING
GO_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION, GO_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION
GO_POSITIVE_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION
 GO_MUSCLE_HYPERTROPHY, GO_MUSCLE_HYPERTROPHY
GO POSITIVE REGULATION OF G1 S TRANSITION OF MITOTIC CELL CYCLE, GO POSITIVE REGULATION OF G1 S TRANSITION OF MITOTIC CELL CYCLE
 GO_REGULATION_OF_VASCULAR_SMOOTH_MUSCLE_CELL_PROLIFERATION, GO_REGULATION_OF_VASCULAR_SMOOTH_MUSCLE_CELL_PROLIFERATION
GO POSITIVE REGULATION OF FATTY ACID METABOLIC PROCESS, GO POSITIVE REGULATION OF FATTY ACID METABOLIC PROCESS
GO_PHENOTYPIC_SWITCHING, GO_PHENOTYPIC_SWITCHING
GO_NEGATIVE_REGULATION_OF_VASCULAR_SMOOTH_MUSCLE_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_VASCULAR_SMOOTH_MUSCLE_CELL_PROLIFERATION.
GO_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS, GO_REGULATION_OF_REACTIVE_OXYGEN_SPECIES_METABOLIC_PROCESS
GO_FORMATION_OF_PRIMARY_GERM_LAYER, GO_FORMATION_OF_PRIMARY_GERM_LAYER
GO_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS, GO_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS
GO_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GO_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING
GO POSITIVE REGULATION OF CARDIAC MUSCLE ADAPTATION, GO POSITIVE REGULATION OF CARDIAC MUSCLE ADAPTATION
 GO_MESODERM_MORPHOGENESIS, GO_MESODERM_MORPHOGENESIS
GO POSITIVE REGULATION OF MICROGLIAL CELL ACTIVATION, GO POSITIVE REGULATION OF MICROGLIAL CELL ACTIVATION
GO_POSITIVE_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS, GO_POSITIVE_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING
 GO_GLIAL_CELL_ACTIVATION, GO_GLIAL_CELL_ACTIVATION
 GO_REGULATION_OF_TISSUE_REMODELING, GO_REGULATION_OF_TISSUE_REMODELING
GO_NEGATIVE_REGULATION_OF_MONOOXYGENASE_ACTIVITY, GO_NEGATIVE_REGULATION_OF_MONOOXYGENASE_ACTIVITY
 GO_POSITIVE_REGULATION_OF_INTERLEUKIN_1_PRODUCTION, GO_POSITIVE_REGULATION_OF_INTERLEUKIN_1_PRODUCTION
GO_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION, GO_REGULATION_OF_CARDIOCYTE_DIFFERENTIATION
GO_REGULATION_OF_SMOOTH_MUSCLE_CONTRACTION, GO_REGULATION_OF_SMOOTH_MUSCLE_CONTRACTION
GO_REACTIVE_NITROGEN_SPECIES_METABOLIC_PROCESS, GO_REACTIVE_NITROGEN_SPECIES_METABOLIC_PROCESS
GO_REGULATION_OF_MACROPHAGE_ACTIVATION, GO_REGULATION_OF_MACROPHAGE_ACTIVATION
GO_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_GROWTH_FACTOR_STIMULUS, GO_NEGATIVE_REGULATION_OF_CELLULAR_RESPONSE_TO_GROWTH_FACTOR_ST
GO_BLOOD_VESSEL_ENDOTHELIAL_CELL_PROLIFERATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS, GO_BLOOD_VESSEL_ENDOTHELIAL_CELL_PROLIFERATION_INVOLVEI
GO_NEGATIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS, GO_NEGATIVE_REGULATION_OF_RECEPTOR_MEDIATED_ENDOCYTOSIS
GO_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL_INVOLVED_IN_CONTRACTION, GO_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL_INVOLVED_IN_CONTRACTION
GO_POSITIVE_REGULATION_OF_STEROID_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_STEROID_METABOLIC_PROCESS
GO_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS, GO_REACTIVE_OXYGEN_SPECIES_BIOSYNTHETIC_PROCESS
GO_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL, GO_CARDIAC_MUSCLE_CELL_ACTION_POTENTIAL
GO_NEGATIVE_REGULATION_OF_CHEMOTAXIS, GO_NEGATIVE_REGULATION_OF_CHEMOTAXIS
GO_NEGATIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS, GO_NEGATIVE_REGULATION_OF_SPROUTING_ANGIOGENESIS
GO_CARDIAC_MUSCLE_CELL_CONTRACTION, GO_CARDIAC_MUSCLE_CELL_CONTRACTION
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_G1_S_PHASE_TRANSITION
GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_MIGRATION
GO_REGULATION_OF_NEUROINFLAMMATORY_RESPONSE, GO_REGULATION_OF_NEUROINFLAMMATORY_RESPONSE
GO_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL, GO_REGULATION_OF_CALCIUM_ION_TRANSPORT_INTO_CYTOSOL
GO_REGULATION_OF_NITRIC_OXIDE_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_NITRIC_OXIDE_BIOSYNTHETIC_PROCESS
GO_MUSCLE_HYPERTROPHY_IN_RESPONSE_TO_STRESS, GO_MUSCLE_HYPERTROPHY_IN_RESPONSE_TO_STRESS
GO_AORTA_MORPHOGENESIS, GO_AORTA_MORPHOGENESIS
GO_GASTRULATION, GO_GASTRULATION
GO_REGULATION_OF_PLASMA_LIPOPROTEIN_PARTICLE_LEVELS, GO_REGULATION_OF_PLASMA_LIPOPROTEIN_PARTICLE_LEVELS
GO REGULATION OF CELL CYCLE G1 S PHASE TRANSITION, GO REGULATION OF CELL CYCLE G1 S PHASE TRANSITION
GO LEUKOCYTE ACTIVATION INVOLVED IN INFLAMMATORY RESPONSE, GO LEUKOCYTE ACTIVATION INVOLVED IN INFLAMMATORY RESPONSE
GO_NEGATIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING, GO_NEGATIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING
GO_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SECRETION, GO_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SECRETION
GO_RESPONSE_TO_STEROL, GO_RESPONSE_TO_STEROL
GO_MEMBRANE_BIOGENESIS, GO_MEMBRANE_BIOGENESIS
GO_CONNECTIVE_TISSUE_REPLACEMENT, GO_CONNECTIVE_TISSUE_REPLACEMENT
GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION, GO_POSITIVE_REGULATION_OF_ENDOTHELIAL_CELL_MIGRATION
GO_NEGATIVE_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_NEGATIVE_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS
GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_CYTOKINE_STIMULUS
GO_NEGATIVE_REGULATION_OF_STEM_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_STEM_CELL_PROLIFERATION
GO_POSITIVE_REGULATION_OF_MUSCLE_ADAPTATION, GO_POSITIVE_REGULATION_OF_MUSCLE_ADAPTATION
GO_POSITIVE_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_ALCOHOL_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_CELLULAR_KETONE_METABOLIC_PROCESS, GO_REGULATION_OF_CELLULAR_KETONE_METABOLIC_PROCESS
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GO_NEGATIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GO_NEGATIVE_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_

GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_CELL_CELL_ADHESION, GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_CELL_CELL_ADHESION

GO_MIRNA_MEDIATED_INHIBITION_OF_TRANSLATION, GO_MIRNA_MEDIATED_INHIBITION_OF_TRANSLATION

GO_REGULATION_OF_RESPONSE_TO_WOUNDING, GO_REGULATION_OF_RESPONSE_TO_WOUNDING

GO_POSITIVE_REGULATION_OF_INFLAMMATORY_RESPONSE, GO_POSITIVE_REGULATION_OF_INFLAMMATORY_RESPONSE