

SIGNALING_PATHWAY,GO_FC_RECEPTOR_SIGNALING_PATHWAY

GO_FICOLIN_1_RICH_GRANULE,GO_FICOLIN_1_RICH_GRANULE
GO_FC_EPSILON_RECEPTOR_SIGNALING_PATHWAY,GO_FC_EPSILON_RECEPTOR_SIGNALING_PATHWAY
GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS,GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
GO_LAMELLIPODIUM,GO_LAMELLIPODIUM
GO_POSITIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY,GO_POSITIVE_REGULATION_OF_NF_KAPPAB_TRANSCRIPTION_FACTOR_ACTIVITY
GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA,GO_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_II_PROMOTER_IN_RESPONSE_TO_HYPOXIA
GO_REGULATION_OF_T_CELL_DIFFERENTIATION,GO_REGULATION_OF_T_CELL_DIFFERENTIATION
GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS,GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS
GO_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM,GO_MORPHOGENESIS_OF_A_POLARIZED_EPITHELIUM
GO_RAS_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY,GO_RAS_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY
GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY,GO_CYTOPLASMIC_PATTERN_RECOGNITION_RECEPTOR_SIGNALING_PATHWAY
GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY,GO_INTERLEUKIN_1_MEDIATED_SIGNALING_PATHWAY
GO_RHO_GTPASE_BINDING,GO_RHO_GTPASE_BINDING
GO_IMMUNOGLOBULIN_BINDING,GO_IMMUNOGLOBULIN_BINDING
GO_PLASMA_MEMBRANE_SIGNALING_RECEPTOR_COMPLEX,GO_PLASMA_MEMBRANE_SIGNALING_RECEPTOR_COMPLEX
GO_LIPID_PHOSPHORYLATION,GO_LIPID_PHOSPHORYLATION
GO_POSITIVE_REGULATION_OF_B_CELL_PROLIFERATION,GO_POSITIVE_REGULATION_OF_B_CELL_PROLIFERATION
GO_PHOSPHATIDYLINOSITOL_PHOSPHORYLATION,GO_PHOSPHATIDYLINOSITOL_PHOSPHORYLATION
GO_PROTEIN_IMPORT,GO_PROTEIN_IMPORT
GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT,GO_POSITIVE_REGULATION_OF_INTRACELLULAR_PROTEIN_TRANSPORT
GO_REGULATION_OF_LIPASE_ACTIVITY,GO_REGULATION_OF_LIPASE_ACTIVITY
GO_HOMOTYPIC_CELL_CELL_ADHESION,GO_HOMOTYPIC_CELL_CELL_ADHESION
GO_TAU_PROTEIN_BINDING,GO_TAU_PROTEIN_BINDING
GO_RUFFLE,GO_RUFFLE
GO_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX,GO_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX
GO_INNATE_IMMUNE_RESPONSE_ACTIVATING_CELL_SURFACE_RECEPTOR_SIGNALING_PATHWAY,GO_INNATE_IMMUNE_RESPONSE_ACTIVATING_CELL_SURFACE_RECEPTOR_SIGNALING_PATHWAY
GO_INTERFERON_GAMMA_MEDIATED_SIGNALING_PATHWAY,GO_INTERFERON_GAMMA_MEDIATED_SIGNALING_PATHWAY
GO_FIBROBLAST_MIGRATION,GO_FIBROBLAST_MIGRATION
GO_EPHRIN_RECEPTOR_SIGNALING_PATHWAY,GO_EPHRIN_RECEPTOR_SIGNALING_PATHWAY
GO_FOCAL_ADHESION_ASSEMBLY,GO_FOCAL_ADHESION_ASSEMBLY
GO_DENDRITIC_SPINE_DEVELOPMENT,GO_DENDRITIC_SPINE_DEVELOPMENT
GO_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION,GO_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION
GO_NEURON_PROJECTION_ORGANIZATION,GO_NEURON_PROJECTION_ORGANIZATION
GO_REGULATION_OF_MYELOID_LEUKOCYTE_MEDIATED_IMMUNITY,GO_REGULATION_OF_MYELOID_LEUKOCYTE_MEDIATED_IMMUNITY
GO_POSITIVE_REGULATION_OF_PHAGOCYTOSIS_ENGULFMENT,GO_POSITIVE_REGULATION_OF_PHAGOCYTOSIS_ENGULFMENT
GO_REGULATION_OF_SUPEROXIDE_ANION_GENERATION,GO_REGULATION_OF_SUPEROXIDE_ANION_GENERATION
GO_DENDRITIC_SPINE_MORPHOGENESIS,GO_DENDRITIC_SPINE_MORPHOGENESIS
GO_RUFFLE_MEMBRANE,GO_RUFFLE_MEMBRANE
GO_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION,GO_POSITIVE_REGULATION_OF_CELL_SUBSTRATE_JUNCTION_ORGANIZATION
GO_NF_KAPPAB_BINDING,GO_NF_KAPPAB_BINDING
GO_SCHAEFFER_COLLATERAL_CA1_SYNAPSE,GO_SCHAEFFER_COLLATERAL_CA1_SYNAPSE
GO_REGULATION_OF_B_CELL_PROLIFERATION,GO_REGULATION_OF_B_CELL_PROLIFERATION
GO_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS,GO_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_JUN_KINASE_ACTIVITY,GO_REGULATION_OF_JUN_KINASE_ACTIVITY
GO_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY,GO_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY
GO_REGULATION_OF_CELL_SIZE,GO_REGULATION_OF_CELL_SIZE
GO_ACTIN_FILAMENT_BASED_MOVEMENT,GO_ACTIN_FILAMENT_BASED_MOVEMENT
GO_PEPTIDYL_THREONINE_MODIFICATION,GO_PEPTIDYL_THREONINE_MODIFICATION
GO_INTERLEUKIN_4_PRODUCTION,GO_INTERLEUKIN_4_PRODUCTION
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER,GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_INITIATION_FROM_RNA_POLYMERASE_II_PROMOTER
GO_MYELOID_LEUKOCYTE_CYTOKINE_PRODUCTION,GO_MYELOID_LEUKOCYTE_CYTOKINE_PRODUCTION
GO_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING,GO_POLYUBIQUITIN_MODIFICATION_DEPENDENT_PROTEIN_BINDING
GO_ACTIN_FILAMENT_BINDING,GO_ACTIN_FILAMENT_BINDING
GO_POSITIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY,GO_POSITIVE_REGULATION_OF_PROTEIN_TYROSINE_KINASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_PRL_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II,GO_POSITIVE_REGULATION_OF_PRL_MIRNA_TRANSCRIPTION_BY_RNA_POLYMERASE_II
GO_REGULATION_OF_INTERLEUKIN_4_PRODUCTION,GO_REGULATION_OF_INTERLEUKIN_4_PRODUCTION
GO_TRANSCRIPTION_COFACTOR_BINDING,GO_TRANSCRIPTION_COFACTOR_BINDING
GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_KINASE_BINDING,GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_KINASE_BINDING
GO_MATURE_B_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE,GO_MATURE_B_CELL_DIFFERENTIATION_INVOLVED_IN_IMMUNE_RESPONSE
GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING,GO_MITOGEN_ACTIVATED_PROTEIN_KINASE_BINDING
GO_REGULATION_OF_CELL_SHAPE,GO_REGULATION_OF_CELL_SHAPE
GO_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT,GO_POSITIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
GO_NEGATIVE_REGULATION_OF_SYNAPSE_ORGANIZATION,GO_NEGATIVE_REGULATION_OF_SYNAPSE_ORGANIZATION
GO_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY,GO_POSITIVE_REGULATION_OF_LAMELLIPODIUM_ASSEMBLY
GO_1_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY,GO_1_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY
GO_REGULATION_OF_NEUTROPHIL_ACTIVATION,GO_REGULATION_OF_NEUTROPHIL_ACTIVATION
GO_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY,GO_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY
GO_POSITIVE_REGULATION_OF_INTERLEUKIN_2_PRODUCTION,GO_POSITIVE_REGULATION_OF_INTERLEUKIN_2_PRODUCTION
GO_1_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATOR_ACTIVITY,GO_1_PHOSPHATIDYLINOSITOL_3_KINASE_REGULATOR_ACTIVITY
GO_POSITIVE_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY,GO_POSITIVE_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY
GO_MYOSIN_COMPLEX,GO_MYOSIN_COMPLEX
GO_POSITIVE_REGULATION_OF_CELL_MATRIX_ADHESION,GO_POSITIVE_REGULATION_OF_CELL_MATRIX_ADHESION
GO_NEURON_PROJECTION_EXTENSION,GO_NEURON_PROJECTION_EXTENSION
GO_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY,GO_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY
GO_AXON_EXTENSION,GO_AXON_EXTENSION
GO_ENDOCYTIC_PATCH,GO_ENDOCYTIC_PATCH
GO_POSITIVE_REGULATION_OF_PROTEIN_IMPORT,GO_POSITIVE_REGULATION_OF_PROTEIN_IMPORT
GO_REGULATION_OF_FIBROBLAST_MIGRATION,GO_REGULATION_OF_FIBROBLAST_MIGRATION
GO_FILOPODIUM,GO_FILOPODIUM