

SIGNALING_PATHWAY, GO_FC_RECEPTOR_SIGNALING_PATHWAY

- GO_LAMELLIPODIUM, GO_LAMELLIPODIUM
- GO_FC_EPSILON_RECEPTOR_SIGNALING_PATHWAY, GO_FC_EPSILON_RECEPTOR_SIGNALING_PATHWAY
- GO_FICOLIN_1_RICH_GRANULE, GO_FICOLIN_1_RICH_GRANULE
- GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION
- 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_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
- GO_POSITIVE_REGULATION_OF_CANONICAL_WNT_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_CANONICAL_WNT_SIGNALING_PATHWAY
- GO_REGULATION_OF_CELL_SHAPE, GO_REGULATION_OF_CELL_SHAPE
- GO_PHAGOCYTIC_VESICLE, GO_PHAGOCYTIC_VESICLE
- GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_DNA_TEMPLATED_TRANSCRIPTION_IN_RESPONSE_TO_STRESS
- GO_FILOPODIUM, GO_FILOPODIUM
- GO GRANULOCYTE MIGRATION, GO GRANULOCYTE MIGRATION
- GO_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION, GO_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION
- GO_POSITIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_WNT_SIGNALING_PATHWAY
- GO_SPECIFIC_GRANULE, GO_SPECIFIC_GRANULE
- GO_RUFFLE, GO_RUFFLE
- GO_POSITIVE_REGULATION_OF_CYTOKINE_PRODUCTION_INVOLVED_IN_IMMUNE_RESPONSE, GO_POSITIVE_REGULATION_OF_CYTOKINE_PRODUCTION_INVOLVED_IN_IMMUNE_RESPONSE
- GO_REGULATION_OF_CELL_KILLING, GO_REGULATION_OF_CELL_KILLING
- GO_ACTIN_FILAMENT_BINDING, GO_ACTIN_FILAMENT_BINDING
- GO_PROTEASOME_ACCESSORY_COMPLEX, GO_PROTEASOME_ACCESSORY_COMPLEX
- GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I, GO_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_I
- GO_POSITIVE_REGULATION_OF_PRODUCTION_OF_MOLECULAR_MEDIATOR_OF_IMMUNE_RESPONSE, GO_POSITIVE_REGULATION_OF_PRODUCTION_OF_MOLECULAR_MEDIATOR_OF_IMMUNE_RESPONSE
- GO_RHO_GTPASE_BINDING, GO_RHO_GTPASE_BINDING
- GO_ACTIN_FILAMENT_POLYMERIZATION, GO_ACTIN_FILAMENT_POLYMERIZATION
- GO_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GO_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
- GO_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY, GO_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY
- GO_RESPONSE_TO_INTERLEUKIN_12, GO_RESPONSE_TO_INTERLEUKIN_12
- GO_PROTEIN_KINASE_C_ACTIVITY, GO_PROTEIN_KINASE_C_ACTIVITY
- GO_POSITIVE_REGULATION_OF_JNK_CASCADE, GO_POSITIVE_REGULATION_OF_JNK_CASCADE
- GO_GROWTH_FACTOR_RECEPTOR_BINDING, GO_GROWTH_FACTOR_RECEPTOR_BINDING
- GO_RESPONSE_TO_FUNGUS, GO_RESPONSE_TO_FUNGUS
- GO_MEMBRANE_INVAGINATION, GO_MEMBRANE_INVAGINATION
- GO_INTERFERON_GAMMA_MEDIATED_SIGNALING_PATHWAY, GO_INTERFERON_GAMMA_MEDIATED_SIGNALING_PATHWAY
- GO_RHO_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY, GO_RHO_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY
- GO_POSITIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION, GO_POSITIVE_REGULATION_OF_CD4_POSITIVE_ALPHA_BETA_T_CELL_ACTIVATION
- GO_POSITIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE, GO_POSITIVE_REGULATION_OF_STRESS_ACTIVATED_PROTEIN_KINASE_SIGNALING_CASCADE
- GO_COP9_SIGNALOSOME, GO_COP9_SIGNALOSOME
- GO_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION, GO_NEGATIVE_REGULATION_OF_B_CELL_ACTIVATION
- GO_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION, GO_REGULATION_OF_HEMATOPOIETIC_PROGENITOR_CELL_DIFFERENTIATION
- GO_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS, GO_REGULATION_OF_CELLULAR_AMINO_ACID_METABOLIC_PROCESS
- GO_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS, GO_PHOSPHATIDYLINOSITOL_BIOSYNTHETIC_PROCESS
- GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_SCF_DEPENDENT_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
- GO_POSITIVE_REGULATION_OF_PROTEIN_POLYMERIZATION, GO_POSITIVE_REGULATION_OF_PROTEIN_POLYMERIZATION
- GO_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION, GO_ARP2_3_COMPLEX_MEDIATED_ACTIN_NUCLEATION
- GO_POSITIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_ALPHA_BETA_T_CELL_DIFFERENTIATION
- GO_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_BINDING, GO_PLATELET_DERIVED_GROWTH_FACTOR_RECEPTOR_BINDING
- GO_POSITIVE_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_PHOSPHOLIPASE_ACTIVITY
- GO_POSITIVE_REGULATION_OF_CYTOKINE_SECRETION, GO_POSITIVE_REGULATION_OF_CYTOKINE_SECRETION
- GO_REGULATION_OF_HOMOTYPIC_CELL_CELL_ADHESION, GO_REGULATION_OF_HOMOTYPIC_CELL_CELL_ADHESION
- GO_REGULATION_OF_PHOSPHOLIPASE_C_ACTIVITY, GO_REGULATION_OF_PHOSPHOLIPASE_C_ACTIVITY
- GO_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GO_INTEGRIN_MEDIATED_SIGNALING_PATHWAY
- GO_REGULATION_OF_T_CELL_RECEPTOR_SIGNALING_PATHWAY, GO_REGULATION_OF_T_CELL_RECEPTOR_SIGNALING_PATHWAY
- GO_RAP_PROTEIN_SIGNAL_TRANSDUCTION, GO_RAP_PROTEIN_SIGNAL_TRANSDUCTION
- GO_T_CELL_PROLIFERATION, GO_T_CELL_PROLIFERATION
- GO_CORTICAL_CYTOSKELETON_ORGANIZATION, GO_CORTICAL_CYTOSKELETON_ORGANIZATION
- GO_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION, GO_HEMATOPOIETIC_STEM_CELL_DIFFERENTIATION
- GO_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY, GO_MYD88_INDEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY
- GO_MAST_CELL_GRANULE, GO_MAST_CELL_GRANULE
- GO_OSTEOCLAST_DIFFERENTIATION, GO_OSTEOCLAST_DIFFERENTIATION
- GO_PHOSPHATIDYLINOSITOL_3_4_5_TRISPHOSPHATE_BINDING, GO_PHOSPHATIDYLINOSITOL_3_4_5_TRISPHOSPHATE_BINDING
- GO_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL
- GO_REGULATION_OF_CELL_JUNCTION_ASSEMBLY, GO_REGULATION_OF_CELL_JUNCTION_ASSEMBLY
- GO_POSITIVE_REGULATION_OF_CYTOKINE_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_CYTOKINE_BIOSYNTHETIC_PROCESS
- GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS, GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS
- GO_REGULATION_OF_MEMBRANE_INVAGINATION, GO_REGULATION_OF_MEMBRANE_INVAGINATION
- GO_CELLULAR_RESPONSE_TO_CADMIUM_ION, GO_CELLULAR_RESPONSE_TO_CADMIUM_ION
- GO_DENDRITIC_SPINE_MORPHOGENESIS, GO_DENDRITIC_SPINE_MORPHOGENESIS
- GO_ACTIN_NUCLEATION, GO_ACTIN_NUCLEATION
- GO_REGULATION_OF_INTERLEUKIN_4_PRODUCTION, GO_REGULATION_OF_INTERLEUKIN_4_PRODUCTION