

TE_MIGRATION, GO_GRANULOCYTE_MIGRATION

- GO_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE, GO_REGULATION_OF_HUMORAL_IMMUNE_RESPONSE
- GO_REGULATION_OF_LEUKOCYTE_MIGRATION, GO_REGULATION_OF_LEUKOCYTE_MIGRATION
- GO_COMPLEMENT_ACTIVATION, GO_COMPLEMENT_ACTIVATION
- GO_CHEMOKINE_RECEPTOR_BINDING, GO_CHEMOKINE_RECEPTOR_BINDING
- GO_CHEMOKINE_ACTIVITY, GO_CHEMOKINE_ACTIVITY
- GO_MONONUCLEAR_CELL_MIGRATION, GO_MONONUCLEAR_CELL_MIGRATION
- GO_INTEGRIN_MEDIATED_SIGNALING_PATHWAY, GO_INTEGRIN_MEDIATED_SIGNALING_PATHWAY
- GO_RESPONSE_TO_CHEMOKINE, GO_RESPONSE_TO_CHEMOKINE
- GO_NEUTROPHIL_MIGRATION, GO_NEUTROPHIL_MIGRATION
- GO_MONOCYTE_CHEMOTAXIS, GO_MONOCYTE_CHEMOTAXIS
- GO_POSITIVE_REGULATION_OF_HEMOPOIESIS, GO_POSITIVE_REGULATION_OF_HEMOPOIESIS
- GO_CELLULAR_DEFENSE_RESPONSE, GO_CELLULAR_DEFENSE_RESPONSE
- GO_CCR_CHEMOKINE_RECEPTOR_BINDING, GO_CCR_CHEMOKINE_RECEPTOR_BINDING
- GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_MIGRATION, GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_MIGRATION
- GO_POSITIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION
- GO_NEGATIVE_REGULATION_OF_CELL_ACTIVATION, GO_NEGATIVE_REGULATION_OF_CELL_ACTIVATION
- GO_POSITIVE_REGULATION_OF_CYTOKINE_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_CYTOKINE_BIOSYNTHETIC_PROCESS
- GO_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_MYELOID_CELL_DIFFERENTIATION
- GO_INTERFERON_GAMMA_PRODUCTION, GO_INTERFERON_GAMMA_PRODUCTION
- GO_OSTEOCLAST_DIFFERENTIATION, GO_OSTEOCLAST_DIFFERENTIATION
- GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION
- GO_MACROPHAGE_MIGRATION, GO_MACROPHAGE_MIGRATION
- GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY, GO_FC_RECEPTOR_MEDIATED_STIMULATORY_SIGNALING_PATHWAY
- GO_POSITIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING, GO_POSITIVE_REGULATION_OF_I_KAPPAB_KINASE_NF_KAPPAB_SIGNALING
- GO_POSITIVE_REGULATION_OF_CELL_KILLING, GO_POSITIVE_REGULATION_OF_CELL_KILLING
- GO_INTERLEUKIN_17_PRODUCTION, GO_INTERLEUKIN_17_PRODUCTION
- GO_PHOSPHOTYROSINE_RESIDUE_BINDING, GO_PHOSPHOTYROSINE_RESIDUE_BINDING
- GO_PHAGOCYTIC_CUP, GO_PHAGOCYTIC_CUP
- GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_10_PRODUCTION, GO_NEGATIVE_REGULATION_OF_INTERLEUKIN_10_PRODUCTION
- GO_T_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE, GO_T_CELL_ACTIVATION_INVOLVED_IN_IMMUNE_RESPONSE
- GO_C_X_C_CHEMOKINE_RECEPTOR_ACTIVITY, GO_C_X_C_CHEMOKINE_RECEPTOR_ACTIVITY
- GO_PHOSPHATIDYLINOSITOL_PHOSPHORYLATION, GO_PHOSPHATIDYLINOSITOL_PHOSPHORYLATION
- GO_MUSCLE_FIBER_DEVELOPMENT, GO_MUSCLE_FIBER_DEVELOPMENT
- GO_ASTROCYTE_DIFFERENTIATION, GO_ASTROCYTE_DIFFERENTIATION
- GO_REGULATION_OF_CYTOLYSIS, GO_REGULATION_OF_CYTOLYSIS
- GO_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE, GO_POSITIVE_REGULATION_OF_ACUTE_INFLAMMATORY_RESPONSE
- GO_REGULATION_OF_MYOBLAST_DIFFERENTIATION, GO_REGULATION_OF_MYOBLAST_DIFFERENTIATION
- GO_NEUTROPHIL_HOMEOSTASIS, GO_NEUTROPHIL_HOMEOSTASIS
- GO_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION, GO_REGULATION_OF_T_CELL_CYTOKINE_PRODUCTION
- GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_CELL_CELL_ADHESION, GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_CELL_CELL_ADHESION
- GO_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION, GO_POSITIVE_REGULATION_OF_INTERLEUKIN_12_PRODUCTION
- GO_B_CELL_PROLIFERATION, GO_B_CELL_PROLIFERATION
- GO_APOPTOTIC_CELL_CLEARANCE, GO_APOPTOTIC_CELL_CLEARANCE
- GO_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY, GO_MYD88_DEPENDENT_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY
- GO_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_ENDOTHELIAL_CELL, GO_POSITIVE_REGULATION_OF_LEUKOCYTE_ADHESION_TO_VASCULAR_E
- GO_GROWTH_FACTOR_RECEPTOR_BINDING, GO_GROWTH_FACTOR_RECEPTOR_BINDING
- GO_POSITIVE_REGULATION_OF_MACROPHAGE_MIGRATION, GO_POSITIVE_REGULATION_OF_MACROPHAGE_MIGRATION
- GO_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION, GO_REGULATION_OF_MONONUCLEAR_CELL_MIGRATION
- GO_LIPID_PHOSPHORYLATION, GO_LIPID_PHOSPHORYLATION
- GO_PROTEIN_COMPLEX_INVOLVED_IN_CELL_ADHESION, GO_PROTEIN_COMPLEX_INVOLVED_IN_CELL_ADHESION