

CELL_HOMEOSTASIS, GO_MYELOID_CELL_HOMEOSTASIS

- GO_PROTEIN_TYROSINE_KINASE_ACTIVITY, GO_PROTEIN_TYROSINE_KINASE_ACTIVITY
- GO_ERYTHROCYTE_HOMEOSTASIS, GO_ERYTHROCYTE_HOMEOSTASIS
- GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_KINASE_ACTIVITY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_KINASE_ACTIVITY
- GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX, GO_TRANSCRIPTIONAL_REPRESSOR_COMPLEX
- GO_TETRAPYRROLE_METABOLIC_PROCESS, GO_TETRAPYRROLE_METABOLIC_PROCESS
- GO_HISTONE_H3_K4_TRIMETHYLATION, GO_HISTONE_H3_K4_TRIMETHYLATION
- GO_LEUKOCYTE_APOPTOTIC_PROCESS, GO_LEUKOCYTE_APOPTOTIC_PROCESS
- GO_NEGATIVE_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_ERYTHROCYTE_DIFFERENTIATION
- GO_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS, GO_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS
- GO_B_CELL_HOMEOSTASIS, GO_B_CELL_HOMEOSTASIS
- GO_B_CELL_APOPTOTIC_PROCESS, GO_B_CELL_APOPTOTIC_PROCESS
- GO_NOTCH_SIGNALING_PATHWAY, GO_NOTCH_SIGNALING_PATHWAY
- GO_GASTRULATION_WITH_MOUTH_FORMING_SECOND, GO_GASTRULATION_WITH_MOUTH_FORMING_SECOND
- GO_INTESTINAL_EPITHELIAL_CELL_DEVELOPMENT, GO_INTESTINAL_EPITHELIAL_CELL_DEVELOPMENT
- GO_MESENCHYMAL_CELL_PROLIFERATION, GO_MESENCHYMAL_CELL_PROLIFERATION
- GO_RESPONSE_TO_INTERLEUKIN_7, GO_RESPONSE_TO_INTERLEUKIN_7
- GO_ANKYRIN_BINDING, GO_ANKYRIN_BINDING
- GO_T_CELL_SELECTION, GO_T_CELL_SELECTION
- GO_INTERLEUKIN_4_PRODUCTION, GO_INTERLEUKIN_4_PRODUCTION
- GO_METANEPHRIC_NEPHRON_MORPHOGENESIS, GO_METANEPHRIC_NEPHRON_MORPHOGENESIS
- GO_T_HELPER_1_CELL_DIFFERENTIATION, GO_T_HELPER_1_CELL_DIFFERENTIATION
- GO_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GO_T_HELPER_1_TYPE_IMMUNE_RESPONSE
- GO_RESPIRATORY_BURST_INVOLVED_IN_DEFENSE_RESPONSE, GO_RESPIRATORY_BURST_INVOLVED_IN_DEFENSE_RESPONSE
- GO_KIDNEY_MORPHOGENESIS, GO_KIDNEY_MORPHOGENESIS
- GO_NEPHRON_MORPHOGENESIS, GO_NEPHRON_MORPHOGENESIS
- GO_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT, GO_CELL_PROLIFERATION_INVOLVED_IN_KIDNEY_DEVELOPMENT
- GO_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION, GO_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION
- GO_OUTFLOW_TRACT_MORPHOGENESIS, GO_OUTFLOW_TRACT_MORPHOGENESIS
- GO_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS, GO_REGULATION_OF_B_CELL_APOPTOTIC_PROCESS
- GO_RENAL_TUBULE_DEVELOPMENT, GO_RENAL_TUBULE_DEVELOPMENT
- GO_POSITIVE_REGULATION_OF_INTERLEUKIN_6_PRODUCTION, GO_POSITIVE_REGULATION_OF_INTERLEUKIN_6_PRODUCTION
- GO_LYSOSOME_LOCALIZATION, GO_LYSOSOME_LOCALIZATION
- GO_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_TO_MESENCHYMAL_TRANSITION
- GO_THYMIC_T_CELL_SELECTION, GO_THYMIC_T_CELL_SELECTION
- GO_CELL_DIFFERENTIATION_INVOLVED_IN_METANEPHROS_DEVELOPMENT, GO_CELL_DIFFERENTIATION_INVOLVED_IN_METANEPHROS_DEVELOPMENT