

NE_ACTIVITY, GO_HORMONE_ACTIVITY

- GO_DRUG_TRANSPORT, GO_DRUG_TRANSPORT
- GO_DIGESTION, GO_DIGESTION
- GO_NEUROPEPTIDE_SIGNALING_PATHWAY, GO_NEUROPEPTIDE_SIGNALING_PATHWAY
- GO_NEUROPEPTIDE_HORMONE_ACTIVITY, GO_NEUROPEPTIDE_HORMONE_ACTIVITY
- GO_GROWTH_FACTOR_ACTIVITY, GO_GROWTH_FACTOR_ACTIVITY
- GO_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION, GO_POSITIVE_REGULATION_OF_SYNAPTIC_TRANSMISSION
- GO_REGULATION_OF_BLOOD_PRESSURE, GO_REGULATION_OF_BLOOD_PRESSURE
- GO_RESPONSE_TO_FOOD, GO_RESPONSE_TO_FOOD
- GO_NEUROPEPTIDE_RECEPTOR_BINDING, GO_NEUROPEPTIDE_RECEPTOR_BINDING
- GO_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT, GO_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT
- GO_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT, GO_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT
- GO_MULTICELLULAR_ORGANISMAL_SIGNALING, GO_MULTICELLULAR_ORGANISMAL_SIGNALING
- GO_GOLGI_LUMEN, GO_GOLGI_LUMEN
- GO_AMINE_TRANSPORT, GO_AMINE_TRANSPORT
- GO_AMMONIUM_TRANSPORT, GO_AMMONIUM_TRANSPORT
- GO_FEEDING_BEHAVIOR, GO_FEEDING_BEHAVIOR
- GO_MONOAMINE_TRANSPORT, GO_MONOAMINE_TRANSPORT
- GO_ACID_SECRETION, GO_ACID_SECRETION
- GO_REGULATION_OF_RESPONSE_TO_FOOD, GO_REGULATION_OF_RESPONSE_TO_FOOD
- GO_NEGATIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_CARTILAGE_DEVELOPMENT
- GO_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION, GO_POSITIVE_REGULATION_OF_BLOOD_CIRCULATION
- GO_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE, GO_REGULATION_OF_SYSTEMIC_ARTERIAL_BLOOD_PRESSURE
- GO_THYROID_HORMONE_METABOLIC_PROCESS, GO_THYROID_HORMONE_METABOLIC_PROCESS
- GO_INOSITOL_LIPID_MEDIATED_SIGNALING, GO_INOSITOL_LIPID_MEDIATED_SIGNALING
- GO_RESPONSE_TO_CORTICOSTEROID, GO_RESPONSE_TO_CORTICOSTEROID
- GO_REGULATION_OF_BEHAVIOR, GO_REGULATION_OF_BEHAVIOR
- GO_INSULIN_RECEPTOR_SIGNALING_PATHWAY, GO_INSULIN_RECEPTOR_SIGNALING_PATHWAY
- GO_BONE_MINERALIZATION, GO_BONE_MINERALIZATION
- GO_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING, GO_PHOSPHATIDYLINOSITOL_3_KINASE_SIGNALING
- GO_SLEEP, GO_SLEEP
- GO_REGULATION_OF_APPETITE, GO_REGULATION_OF_APPETITE
- GO_NEGATIVE_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS, GO_NEGATIVE_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS
- GO_ADENYLATE_CYCLASE_ACTIVATING_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GO_ADENYLATE_CYCLASE_ACTIVATING_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY
- GO_PHOSPHOLIPASE_C_ACTIVATING_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GO_PHOSPHOLIPASE_C_ACTIVATING_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY
- GO_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GO_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
- GO_REGULATION_OF_SENSORY_PERCEPTION, GO_REGULATION_OF_SENSORY_PERCEPTION
- GO_POSITIVE_REGULATION_OF_CALCIIUM_ION_IMPORT, GO_POSITIVE_REGULATION_OF_CALCIIUM_ION_IMPORT
- GO_REGULATION_OF_INTESTINAL_ABSORPTION, GO_REGULATION_OF_INTESTINAL_ABSORPTION
- GO_MACROPHAGE_DIFFERENTIATION, GO_MACROPHAGE_DIFFERENTIATION
- GO_GASTRO_INTESTINAL_SYSTEM_SMOOTH_MUSCLE_CONTRACTION, GO_GASTRO_INTESTINAL_SYSTEM_SMOOTH_MUSCLE_CONTRACTION
- GO_HORMONE_RECEPTOR_BINDING, GO_HORMONE_RECEPTOR_BINDING
- GO_MAGNESIUM_ION_HOMEOSTASIS, GO_MAGNESIUM_ION_HOMEOSTASIS
- GO_RESPONSE_TO_STARVATION, GO_RESPONSE_TO_STARVATION
- GO_MONOSACCHARIDE_BIOSYNTHETIC_PROCESS, GO_MONOSACCHARIDE_BIOSYNTHETIC_PROCESS
- GO_CATECHOLAMINE_SECRETION, GO_CATECHOLAMINE_SECRETION
- GO_ENDOSOME_LUMEN, GO_ENDOSOME_LUMEN
- GO_SENSORY_PERCEPTION_OF_PAIN, GO_SENSORY_PERCEPTION_OF_PAIN
- GO_NOREPINEPHRINE_TRANSPORT, GO_NOREPINEPHRINE_TRANSPORT
- GO_NEGATIVE_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_CHONDROCYTE_DIFFERENTIATION
- GO_GLUCCORTICOID_METABOLIC_PROCESS, GO_GLUCCORTICOID_METABOLIC_PROCESS
- GO_TRANSMISSION_OF_NERVE_IMPULSE, GO_TRANSMISSION_OF_NERVE_IMPULSE
- GO_DIGESTIVE_SYSTEM_PROCESS, GO_DIGESTIVE_SYSTEM_PROCESS
- GO_ASSOCIATIVE_LEARNING, GO_ASSOCIATIVE_LEARNING
- GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
- GO_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS, GO_REGULATION_OF_LIPOPROTEIN_METABOLIC_PROCESS
- GO_REGULATION_OF_HEART_RATE, GO_REGULATION_OF_HEART_RATE
- GO_EXCRETION, GO_EXCRETION
- GO_ACUTE_PHASE_RESPONSE, GO_ACUTE_PHASE_RESPONSE
- GO_VASCULAR_PROCESS_IN_CIRCULATORY_SYSTEM, GO_VASCULAR_PROCESS_IN_CIRCULATORY_SYSTEM
- GO_REGULATION_OF_VASCULAR_PERMEABILITY, GO_REGULATION_OF_VASCULAR_PERMEABILITY
- GO_MAIN_AXON, GO_MAIN_AXON
- GO_REGULATION_OF_CARTILAGE_DEVELOPMENT, GO_REGULATION_OF_CARTILAGE_DEVELOPMENT
- GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_LEUKOCYTE_DIFFERENTIATION
- GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_EXTRACELLULAR_STIMULUS, GO_NEGATIVE_REGULATION_OF_RESPONSE_TO_EXTRACELLULAR_STIMULUS
- GO_RESPONSE_TO_ELECTRICAL_STIMULUS, GO_RESPONSE_TO_ELECTRICAL_STIMULUS
- GO_EXPLORATION_BEHAVIOR, GO_EXPLORATION_BEHAVIOR
- GO_POSITIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT, GO_POSITIVE_REGULATION_OF_RECEPTOR_SIGNALING_PATHWAY_VIA_STAT
- GO_ENDOCRINE_SYSTEM_DEVELOPMENT, GO_ENDOCRINE_SYSTEM_DEVELOPMENT
- GO_POSITIVE_REGULATION_OF_T_CELL_PROLIFERATION, GO_POSITIVE_REGULATION_OF_T_CELL_PROLIFERATION
- GO_NEGATIVE_REGULATION_OF_NOREPINEPHRINE_SECRETION, GO_NEGATIVE_REGULATION_OF_NOREPINEPHRINE_SECRETION
- GO_STARTLE_RESPONSE, GO_STARTLE_RESPONSE
- GO_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN, GO_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN
- GO_POSITIVE_REGULATION_OF_CIRCADIAN_RHYTHM, GO_POSITIVE_REGULATION_OF_CIRCADIAN_RHYTHM
- GO_RESPONSE_TO_MINERALOCORTICOID, GO_RESPONSE_TO_MINERALOCORTICOID
- GO_REGULATION_OF_FEEDING_BEHAVIOR, GO_REGULATION_OF_FEEDING_BEHAVIOR
- GO_PLACENTA_DEVELOPMENT, GO_PLACENTA_DEVELOPMENT
- GO_NEUTROPHIL_MIGRATION, GO_NEUTROPHIL_MIGRATION
- GO_POSITIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN, GO_POSITIVE_REGULATION_OF_TYROSINE_PHOSPHORYLATION_OF_STAT_PROTEIN
- GO_REGULATION_OF_CALCIIUM_ION_IMPORT, GO_REGULATION_OF_CALCIIUM_ION_IMPORT
- GO_FEAR_RESPONSE, GO_FEAR_RESPONSE
- GO_POSITIVE_REGULATION_OF_REPRODUCTIVE_PROCESS, GO_POSITIVE_REGULATION_OF_REPRODUCTIVE_PROCESS
- GO_NEGATIVE_REGULATION_OF_BEHAVIOR, GO_NEGATIVE_REGULATION_OF_BEHAVIOR
- GO_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION, GO_REGULATION_OF_MYELOID_LEUKOCYTE_DIFFERENTIATION
- GO_ACUTE_INFLAMMATORY_RESPONSE_TO_ANTIGENIC_STIMULUS, GO_ACUTE_INFLAMMATORY_RESPONSE_TO_ANTIGENIC_STIMULUS
- GO_PHASIC_SMOOTH_MUSCLE_CONTRACTION, GO_PHASIC_SMOOTH_MUSCLE_CONTRACTION
- GO_CYCLIC_NUCLEOTIDE_BIOSYNTHETIC_PROCESS, GO_CYCLIC_NUCLEOTIDE_BIOSYNTHETIC_PROCESS
- GO_PITUITARY_GLAND_DEVELOPMENT, GO_PITUITARY_GLAND_DEVELOPMENT
- GO_ADULT_HEART_DEVELOPMENT, GO_ADULT_HEART_DEVELOPMENT
- GO_OPIOID_RECEPTOR_BINDING, GO_OPIOID_RECEPTOR_BINDING
- GO_NEGATIVE_REGULATION_OF_AMINE_TRANSPORT, GO_NEGATIVE_REGULATION_OF_AMINE_TRANSPORT
- GO_ACUTE_INFLAMMATORY_RESPONSE, GO_ACUTE_INFLAMMATORY_RESPONSE
- GO_POSITIVE_REGULATION_OF_HEART_CONTRACTION, GO_POSITIVE_REGULATION_OF_HEART_CONTRACTION
- GO_REGULATION_OF_RESPONSE_TO_EXTRACELLULAR_STIMULUS, GO_REGULATION_OF_RESPONSE_TO_EXTRACELLULAR_STIMULUS
- GO_DIGESTIVE_SYSTEM_DEVELOPMENT, GO_DIGESTIVE_SYSTEM_DEVELOPMENT
- GO_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS, GO_REGULATION_OF_DIGESTIVE_SYSTEM_PROCESS
- GO_LEUKOCYTE_TETHERING_OR_ROLLING, GO_LEUKOCYTE_TETHERING_OR_ROLLING
- GO_ACYLGLYCEROL_HOMEOSTASIS, GO_ACYLGLYCEROL_HOMEOSTASIS
- GO_BROWN_FAT_CELL_DIFFERENTIATION, GO_BROWN_FAT_CELL_DIFFERENTIATION
- GO_POLYSACCHARIDE_METABOLIC_PROCESS, GO_POLYSACCHARIDE_METABOLIC_PROCESS
- GO_PEPTIDE_HORMONE_PROCESSING, GO_PEPTIDE_HORMONE_PROCESSING
- GO_FEMALE_SEX_DIFFERENTIATION, GO_FEMALE_SEX_DIFFERENTIATION
- GO_MULTICELLULAR_ORGANISMAL_RESPONSE_TO_STRESS, GO_MULTICELLULAR_ORGANISMAL_RESPONSE_TO_STRESS
- GO_RESPONSE_TO_CHEMOKINE, GO_RESPONSE_TO_CHEMOKINE
- GO_RESPONSE_TO_CORTICOSTERONE, GO_RESPONSE_TO_CORTICOSTERONE