

GOBP_COLLAGEN_ACTIVATED_TYROSINE_KINASE_RECEPTOR_SIGNALING_PATHWAY-

GOBP_COLLAGEN_ACTIVATED_SIGNALING_PATHWAY-

GOBP_CELL_CELL_ADHESION_MEDIATED_BY_INTEGRIN-

GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_PEPTIDE_OR_POLYSACCHARIDE_ANTIGEN_VIA_MHC_CLASS_II-

GOBP_ANTIGEN_PROCESSING_AND_PRESENTATION_OF_EXOGENOUS_PEPTIDE_ANTIGEN_VIA_MHC_CLASS_II-GOBP_DENDRITIC_CELL_ANTIGEN_PROCESSING_AND_PRESENTATION-

GOBP_INFLAMMATORY_RESPONSE_TO_WOUNDING-

GOBP_LYMPH_VESSEL_DEVELOPMENT-

GOBP_CHEMICAL_HOMEOSTASIS_WITHIN_A_TISSUE-

GOBP_NEGATIVE_REGULATION_OF_FATTY_ACID_BIOSYNTHETIC_PROCESS-

GOBP_MEMBRANE_RAFT_ORGANIZATION-

GOBP_RENAL_FILTRATION-

GOBP_MEMBRANE_RAFT_ASSEMBLY-

GOBP_BASEMENT_MEMBRANE_ORGANIZATION-GOBP_MORPHOGENESIS_OF_AN_ENDOTHELIUM-

GOBP_MACROPHAGE_CYTOKINE_PRODUCTION-

GOBP_ELASTIC_FIBER_ASSEMBLY-

GOBP_ASCENDING_AORTA_DEVELOPMENT-

GOBP_NEGATIVE_REGULATION_OF_CELL_MIGRATION_INVOLVED_IN_SPROUTING_ANGIOGENESIS-

GOBP_DICHOTOMOUS_SUBDIVISION_OF_AN_EPITHELIAL_TERMINAL_UNIT-

GOBP_PLASMA_MEMBRANE_RAFT_ASSEMBLY-

GOBP_CAVEOLA_ASSEMBLY-GOBP_NATURAL_KILLER_CELL_CHEMOTAXIS-

GOBP_COMMON_PARTNER_SMAD_PROTEIN_PHOSPHORYLATION-

GOBP_TRIGLYCERIDE_RICH_LIPOPROTEIN_PARTICLE_REMODELING-

GOBP_FATTY_ACID_BETA_OXIDATION_USING_ACYL_COA_OXIDASE-

GOBP_AROMATIC_AMINO_ACID_FAMILY_METABOLIC_PROCESS-

GOBP_TYROSINE_METABOLIC_PROCESS-

GOBP_AROMATIC_AMINO_ACID_FAMILY_CATABOLIC_PROCESS

GOBP_BENZENE_CONTAINING_COMPOUND_METABOLIC_PROCESS-

GOBP_SERINE_FAMILY_AMINO_ACID_CATABOLIC_PROCESS-

GOBP_BRANCHED_CHAIN_AMINO_ACID_CATABOLIC_PROCESS-

GOBP_AMINO_ACID_BETAINE_METABOLIC_PROCESS-

GOBP_BILE_ACID_AND_BILE_SALT_TRANSPORT-

GOBP_REGULATION_OF_LIPOPROTEIN_LIPASE_ACTIVITY-

GOBP_POSITIVE_REGULATION_OF_TRIGLYCERIDE_METABOLIC_PROCESS-

GOBP_DOPAMINE_SECRETION-

GOBP_CATECHOLAMINE_SECRETION-

GOBP_LONG_TERM_SYNAPTIC_POTENTIATION-

GOBP_TRANSMISSION_OF_NERVE_IMPULSE-GOBP_PROTEIN_LOCALIZATION_TO_SYNAPSE-

GOBP_GLUTAMATE_RECEPTOR_SIGNALING_PATHWAY-

GOBP_MODULATION_OF_EXCITATORY_POSTSYNAPTIC_POTENTIAL-

GOBP_CALCIUM_ION_REGULATED_EXOCYTOSIS_OF_NEUROTRANSMITTER-

GOBP_RECEPTOR_LOCALIZATION_TO_SYNAPSE-

GOBP_SYNAPTIC_VESICLE_PRIMING

GOBP_WALKING_BEHAVIOR-GOBP_NEURON_CELL_CELL_ADHESION-

GOBP_HUMORAL_IMMUNE_RESPONSE_MEDIATED_BY_CIRCULATING_IMMUNOGLOBULIN-

GOBP_BILE_ACID_BIOSYNTHETIC_PROCESS-

GOBP_SYNAPTIC_VESICLE_RECYCLING-

GOBP_PRESYNAPTIC_ENDOCYTOSIS-GOBP_REGULATION_OF_NEURONAL_SYNAPTIC_PLASTICITY-

GOBP_REGULATION_OF_SYNAPTIC_VESICLE_EXOCYTOSIS-

GOBP_RETROGRADE_AXONAL_TRANSPORT-

GOBP_ANTEROGRADE_AXONAL_TRANSPORT-

GOBP_ESTABLISHMENT_OF_MITOCHONDRION_LOCALIZATION-

GOBP_AXONAL_TRANSPORT_OF_MITOCHONDRIONGOBP_AXONAL_TRANSPORT-

GOBP_AXO_DENDRITIC_TRANSPORT-

GOBP_LONG_TERM_SYNAPTIC_DEPRESSION-

GOBP_SYNAPTIC_TRANSMISSION_GABAERGIC-GOBP_REGULATION_OF_POSTSYNAPTIC_NEUROTRANSMITTER_RECEPTOR_ACTIVITY-

GOBP_REGULATION_OF_NEUROTRANSMITTER_RECEPTOR_ACTIVITY-GOBP_REGULATION_OF_SYNAPTIC_TRANSMISSION_GLUTAMATERGIC-

GOBP_POSITIVE_REGULATION_OF_SYNAPSE_ASSEMBLY-

GOBP_SERINE_FAMILY_AMINO_ACID_METABOLIC_PROCESS-

GOBP_UREA_CYCLE-

GOBP_UREA_METABOLIC_PROCESS-

GOBP_INTRACILIARY_RETROGRADE_TRANSPORT-