

IAL_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_PROTEIN_CATABOLIC_PROCESS

GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_REGULATION_OF_PROTEASOMAL_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS, KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS
GO_UBIQUITIN_LIKE_PROTEIN_LIGASE_ACTIVITY, GO_UBIQUITIN_LIKE_PROTEIN_LIGASE_ACTIVITY
GSE45365_HEALTHY_VS_MCMV_INFECTION_BCELL_IFNAR_KO_UP, GSE45365_HEALTHY_VS_MCMV_INFECTION_BCELL_IFNAR_KO_UP
GO_PROTEIN_UBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS, GO_PROTEIN_UBIQUITINATION_INVOLVED_IN_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS
GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN, GO_CELLULAR_RESPONSE_TO_TOPOLOGICALLY_INCORRECT_PROTEIN
SMIRNOV_RESPONSE_TO_IR_6HR_UP, SMIRNOV_RESPONSE_TO_IR_6HR_UP
GO_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX, GO_CULLIN_RING_UBIQUITIN_LIGASE_COMPLEX
REACTOME_RECRUITMENT_OF_MITOTIC_CENTROSOME_PROTEINS_AND_COMPLEXES, REACTOME_RECRUITMENT_OF_MITOTIC_CENTROSOME_PROTEINS_AND_COMPLEXES
GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_UP, GSE18804_SPLEEN_MACROPHAGE_VS_TUMORAL_MACROPHAGE_UP
GO_REGULATION_OF_CYTOKINESIS, GO_REGULATION_OF_CYTOKINESIS
NAKAJIMA_MAST_CELL, NAKAJIMA_MAST_CELL
GO_NEGATIVE_REGULATION_OF_CELL_ACTIVATION, GO_NEGATIVE_REGULATION_OF_CELL_ACTIVATION
GO_REGULATION_OF_TRANSPORTER_ACTIVITY, GO_REGULATION_OF_TRANSPORTER_ACTIVITY
GO_REGULATION_OF_NEUROTRANSMITTER_LEVELS, GO_REGULATION_OF_NEUROTRANSMITTER_LEVELS
GO_KINASE_INHIBITOR_ACTIVITY, GO_KINASE_INHIBITOR_ACTIVITY
GO_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
KERLEY_RESPONSE_TO_CISPLATIN_UP, KERLEY_RESPONSE_TO_CISPLATIN_UP
REACTOME_SIGNALING_BY_PDGF, REACTOME_SIGNALING_BY_PDGF
KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION, KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION
GO_POSITIVE_REGULATION_OF_REPRODUCTIVE_PROCESS, GO_POSITIVE_REGULATION_OF_REPRODUCTIVE_PROCESS
GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION, GO_POSITIVE_REGULATION_OF_CHROMOSOME_SEGREGATION
GO_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION, GO_REGULATION_OF_STRIATED_MUSCLE_CONTRACTION
GO_REGULATION_OF_HEART_RATE, GO_REGULATION_OF_HEART_RATE
GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_16H_DN, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_16H_DN
GO_REGULATION_OF_CATION_CHANNEL_ACTIVITY, GO_REGULATION_OF_CATION_CHANNEL_ACTIVITY
GO_POTASSIUM_CHANNEL_REGULATOR_ACTIVITY, GO_POTASSIUM_CHANNEL_REGULATOR_ACTIVITY
LI_PROSTATE_CANCER_EPIGENETIC, LI_PROSTATE_CANCER_EPIGENETIC
GO_CELLULAR_SENESCENCE, GO_CELLULAR_SENESCENCE
GO_GROWTH_FACTOR_RECEPTOR_BINDING, GO_GROWTH_FACTOR_RECEPTOR_BINDING
GO_STARTLE_RESPONSE, GO_STARTLE_RESPONSE
GO_DEVELOPMENTAL_MATURATION, GO_DEVELOPMENTAL_MATURATION
GO_PRESYNAPTIC_PROCESS_INVOLVED_IN_SYNAPTIC_TRANSMISSION, GO_PRESYNAPTIC_PROCESS_INVOLVED_IN_SYNAPTIC_TRANSMISSION
GO_STRIATED_MUSCLE_CELL_DIFFERENTIATION, GO_STRIATED_MUSCLE_CELL_DIFFERENTIATION
GO_REGULATION_OF_DENDRITE_MORPHOGENESIS, GO_REGULATION_OF_DENDRITE_MORPHOGENESIS
GO_FRIZZLED_BINDING, GO_FRIZZLED_BINDING
GO_GLAND_MORPHOGENESIS, GO_GLAND_MORPHOGENESIS
GO_DOPAMINE_TRANSPORT, GO_DOPAMINE_TRANSPORT
GO_RESPONSE_TO_EPINEPHRINE, GO_RESPONSE_TO_EPINEPHRINE
GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT, GO_REGULATION_OF_POTASSIUM_ION_TRANSMEMBRANE_TRANSPORT
GO_14_3_3_PROTEIN_BINDING, GO_14_3_3_PROTEIN_BINDING
HOFMANN_CELL_LYMPHOMA_DN, HOFMANN_CELL_LYMPHOMA_DN
GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY
GO_MONOAMINE_TRANSPORT, GO_MONOAMINE_TRANSPORT
KEGG_BASAL_CELL_CARCINOMA, KEGG_BASAL_CELL_CARCINOMA
GO_CELL_SURFACE_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_CELL_CELL_SIGNALING, GO_CELL_SURFACE_RECEPTOR_SIGNALING_PATHWAY_INVOLVED_IN_CELL_CELL_SIGNALING
GO_REGULATION_OF_DOPAMINE_SECRETION, GO_REGULATION_OF_DOPAMINE_SECRETION
PUJANA_BREAST_CANCER_WITH_BRCA1_MUTATED_DN, PUJANA_BREAST_CANCER_WITH_BRCA1_MUTATED_DN
GO_DERLIN_1_RETROTRANSLOCATION_COMPLEX, GO_DERLIN_1_RETROTRANSLOCATION_COMPLEX
GO_REGULATION_OF_EMBRYONIC_DEVELOPMENT, GO_REGULATION_OF_EMBRYONIC_DEVELOPMENT
GO_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE, GO_REGULATION_OF_T_HELPER_1_TYPE_IMMUNE_RESPONSE
ST_JAK_STAT_PATHWAY, ST_JAK_STAT_PATHWAY
GO_REGULATION_OF_MEIOTIC_CELL_CYCLE, GO_REGULATION_OF_MEIOTIC_CELL_CYCLE
GO_TAU_PROTEIN_KINASE_ACTIVITY, GO_TAU_PROTEIN_KINASE_ACTIVITY