COMPLEX_DISASSEMBLY, GO_MACROMOLECULAR_COMPLEX_DISASSEMBLY

GO_SISTER_CHROMATID_SEGREGATION, GO_SISTER_CHROMATID_SEGREGATION
GSE19825_NAIVE_VS_IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN, GSE19825_NAIVE_VS_IL2RAHIGH_DAY3_EFF_CD8_TCELL_DN
GSE17721_POLYIC_VS_CPG_4H_BMDC_UP, GSE17721_POLYIC_VS_CPG_4H_BMDC_UP
GO_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION, GO_REGULATION_OF_MICROTUBULE_POLYMERIZATION_OR_DEPOLYMERIZATION
GGAANCGGAANY_UNKNOWN, GGAANCGGAANY_UNKNOWN
GSE17721_PAM3CSK4_VS_CPG_8H_BMDC_UP, GSE17721_PAM3CSK4_VS_CPG_8H_BMDC_UP
GO_MITOTIC_SISTER_CHROMATID_SEGREGATION, GO_MITOTIC_SISTER_CHROMATID_SEGREGATION
GO_CHROMATIN_REMODELING, GO_CHROMATIN_REMODELING
GO_REGULATION_OF_NUCLEAR_DIVISION, GO_REGULATION_OF_NUCLEAR_DIVISION
KEGG_ENDOCYTOSIS, KEGG_ENDOCYTOSIS
GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY, GO_CHROMATIN_ASSEMBLY_OR_DISASSEMBLY
GSE3982_DC_VS_TH2_DN, GSE3982_DC_VS_TH2_DN
GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS, GO_POSITIVE_REGULATION_OF_VIRAL_PROCESS
BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_CIS, BYSTRYKH_HEMATOPOIESIS_STEM_CELL_AND_BRAIN_QTL_CIS
GO_ATP_DEPENDENT_CHROMATIN_REMODELING, GO_ATP_DEPENDENT_CHROMATIN_REMODELING
GSE45365_CD8A_DC_VS_CD11B_DC_UP, GSE45365_CD8A_DC_VS_CD11B_DC_UP
GO_MICROTUBULE_ASSOCIATED_COMPLEX, GO_MICROTUBULE_ASSOCIATED_COMPLEX
GO_RIBOSOME_BINDING, GO_RIBOSOME_BINDING
GO_CHROMOSOME_LOCALIZATION, GO_CHROMOSOME_LOCALIZATION
GO_REGULATION_OF_PROTEIN_BINDING, GO_REGULATION_OF_PROTEIN_BINDING
GO_METAPHASE_PLATE_CONGRESSION, GO_METAPHASE_PLATE_CONGRESSION
GO_SITE_OF_POLARIZED_GROWTH, GO_SITE_OF_POLARIZED_GROWTH
GO_CYTOKINESIS, GO_CYTOKINESIS
TNCATNTCCYR_UNKNOWN, TNCATNTCCYR_UNKNOWN
KEGG_WNT_SIGNALING_PATHWAY, KEGG_WNT_SIGNALING_PATHWAY
GO_BAF_TYPE_COMPLEX, GO_BAF_TYPE_COMPLEX
GO_REGULATION_OF_EMBRYONIC_DEVELOPMENT, GO_REGULATION_OF_EMBRYONIC_DEVELOPMENT
GSE21033_CTRL_VS_POLYIC_STIM_DC_6H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_6H_DN
KEGG_COLORECTAL_CANCER, KEGG_COLORECTAL_CANCER
GO_ATP_DEPENDENT_MICROTUBULE_MOTOR_ACTIVITY, GO_ATP_DEPENDENT_MICROTUBULE_MOTOR_ACTIVITY
GO_NBAF_COMPLEX, GO_NBAF_COMPLEX
GO_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_JUN_KINASE_ACTIVITY
GO_DENDRITIC_SHAFT, GO_DENDRITIC_SHAFT
GO_CARDIAC_CHAMBER_DEVELOPMENT, GO_CARDIAC_CHAMBER_DEVELOPMENT
GO_MULTI_ORGANISM_MEMBRANE_ORGANIZATION, GO_MULTI_ORGANISM_MEMBRANE_ORGANIZATION
GO_OUTFLOW_TRACT_MORPHOGENESIS, GO_OUTFLOW_TRACT_MORPHOGENESIS
GO_STEM_CELL_DIFFERENTIATION, GO_STEM_CELL_DIFFERENTIATION
GO_RAC_GTPASE_BINDING, GO_RAC_GTPASE_BINDING
GO_DEVELOPMENTAL_INDUCTION, GO_DEVELOPMENTAL_INDUCTION
GO_SYNAPSE_ORGANIZATION, GO_SYNAPSE_ORGANIZATION
GSE17721_CTRL_VS_POLYIC_0.5H_BMDC_DN, GSE17721_CTRL_VS_POLYIC_0.5H_BMDC_DN
GO_MESENCHYMAL_CELL_DIFFERENTIATION, GO_MESENCHYMAL_CELL_DIFFERENTIATION
REACTOME_FACTORS_INVOLVED_IN_MEGAKARYOCYTE_DEVELOPMENT_AND_PLATELET_PRODUCTION, REACTOME_FACTORS_INVOLVED_IN_MEGAKARYOCYTE_DEVELOPMENT_AND_PLATELET_PRODUCTION
NAKAMURA_ADIPOGENESIS_LATE_UP, NAKAMURA_ADIPOGENESIS_LATE_UP
GSE41176_WT_VS_TAK1_KO_ANTI_IGM_STIM_BCELL_3H_UP, GSE41176_WT_VS_TAK1_KO_ANTI_IGM_STIM_BCELL_3H_UP
GO_GASTRULATION, GO_GASTRULATION
GO_KINESIN_COMPLEX, GO_KINESIN_COMPLEX
GO_NEURAL_TUBE_FORMATION, GO_NEURAL_TUBE_FORMATION
GO_REGULATION_OF_DENDRITE_DEVELOPMENT, GO_REGULATION_OF_DENDRITE_DEVELOPMENT
GO_SPINDLE_MIDZONE, GO_SPINDLE_MIDZONE
GO_ECTODERMAL_PLACODE_DEVELOPMENT, GO_ECTODERMAL_PLACODE_DEVELOPMENT
GO_STRIATED_MUSCLE_CELL_DIFFERENTIATION, GO_STRIATED_MUSCLE_CELL_DIFFERENTIATION
PID_WNT_CANONICAL_PATHWAY, PID_WNT_CANONICAL_PATHWAY
GO_POSITIVE_REGULATION_OF_EMBRYONIC_DEVELOPMENT, GO_POSITIVE_REGULATION_OF_EMBRYONIC_DEVELOPMENT
GO_ACTIN_FILAMENT_ORGANIZATION, GO_ACTIN_FILAMENT_ORGANIZATION
GO_MICROTUBULE_MOTOR_ACTIVITY, GO_MICROTUBULE_MOTOR_ACTIVITY
PID_BETA_CATENIN_NUC_PATHWAY, PID_BETA_CATENIN_NUC_PATHWAY
BIOCARTA_PS1_PATHWAY, BIOCARTA_PS1_PATHWAY
GO_SYNAPSE_ASSEMBLY, GO_SYNAPSE_ASSEMBLY
MATTHEWS_AP1_TARGETS, MATTHEWS_AP1_TARGETS
GO_SWI_SNF_COMPLEX, GO_SWI_SNF_COMPLEX
BROWNE_HCMV_INFECTION_8HR_DN, BROWNE_HCMV_INFECTION_8HR_DN
PID_WNT_SIGNALING_PATHWAY, PID_WNT_SIGNALING_PATHWAY
GO_REGULATION_OF_COLLATERAL_SPROUTING, GO_REGULATION_OF_COLLATERAL_SPROUTING
GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_BODY_SORTING_PATHWAY, GO_UBIQUITIN_DEPENDENT_PROTEIN_CATABOLIC_PROCESS_VIA_THE_MULTIVESICULAR_B
GO_DORSAL_VENTRAL_AXIS_SPECIFICATION, GO_DORSAL_VENTRAL_AXIS_SPECIFICATION
GO_CARDIAC_CHAMBER_MORPHOGENESIS, GO_CARDIAC_CHAMBER_MORPHOGENESIS
GO_MULTIVESICULAR_BODY_ORGANIZATION, GO_MULTIVESICULAR_BODY_ORGANIZATION
GO_MULTI_ORGANISM_ORGANELLE_ORGANIZATION, GO_MULTI_ORGANISM_ORGANELLE_ORGANIZATION
GO_NEGATIVE_REGULATION_OF_PROTEIN_BINDING, GO_NEGATIVE_REGULATION_OF_PROTEIN_BINDING
GO_ESCRT_COMPLEX, GO_ESCRT_COMPLEX
GO_RECEPTOR_REGULATOR_ACTIVITY, GO_RECEPTOR_REGULATOR_ACTIVITY
GO_EYE_MORPHOGENESIS, GO_EYE_MORPHOGENESIS
GO_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS, GO_REGULATION_OF_MUSCLE_CELL_APOPTOTIC_PROCESS
GO_REGULATION_OF_JUN_KINASE_ACTIVITY, GO_REGULATION_OF_JUN_KINASE_ACTIVITY
NADLER_HYPERGLYCEMIA_AT_OBESITY, NADLER_HYPERGLYCEMIA_AT_OBESITY
GO_CARDIAC_SEPTUM_MORPHOGENESIS, GO_CARDIAC_SEPTUM_MORPHOGENESIS
GO_INNER_EAR_MORPHOGENESIS, GO_INNER_EAR_MORPHOGENESIS
LI_PROSTATE_CANCER_EPIGENETIC, LI_PROSTATE_CANCER_EPIGENETIC
GO_EMBRYONIC_CAMERA_TYPE_EYE_MORPHOGENESIS, GO_EMBRYONIC_CAMERA_TYPE_EYE_MORPHOGENESIS
GO_EMBRYONIC_EYE_MORPHOGENESIS, GO_EMBRYONIC_EYE_MORPHOGENESIS
GO_POSITIVE_REGULATION_OF_AXONOGENESIS, GO_POSITIVE_REGULATION_OF_AXONOGENESIS
GO_POSITIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION, GO_POSITIVE_REGULATION_OF_CELL_CYCLE_G2_M_PHASE_TRANSITION
GO_SYNAPTIC_VESICLE_RECYCLING, GO_SYNAPTIC_VESICLE_RECYCLING
GO_POSITIVE_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL, GO_POSITIVE_REGULATION_OF_VIRAL_RELEASE_FROM_HOST_CELL
GO_EMBRYONIC_SKELETAL_JOINT_DEVELOPMENT, GO_EMBRYONIC_SKELETAL_JOINT_DEVELOPMENT
GO_REGULATION_OF_FAT_CELL_DIFFERENTIATION, GO_REGULATION_OF_FAT_CELL_DIFFERENTIATION
GO_FOREBRAIN_REGIONALIZATION, GO_FOREBRAIN_REGIONALIZATION
GO_POSITIVE_REGULATION_OF_DENDRITE_DEVELOPMENT, GO_POSITIVE_REGULATION_OF_DENDRITE_DEVELOPMENT
GO_LATERAL_PLASMA_MEMBRANE, GO_LATERAL_PLASMA_MEMBRANE
PTEN_DN.V2_DN, PTEN_DN.V2_DN
GO_RECEPTOR_ACTIVATOR_ACTIVITY, GO_RECEPTOR_ACTIVATOR_ACTIVITY
GO_CHROMATIN_DISASSEMBLY, GO_CHROMATIN_DISASSEMBLY
GO_RECEPTOR_AGONIST_ACTIVITY, GO_RECEPTOR_AGONIST_ACTIVITY
GO_MORPHOGENESIS_OF_AN_EPITHELIAL_SHEET, GO_MORPHOGENESIS_OF_AN_EPITHELIAL_SHEET