GO_MIDBODY, GO_MIDBODY GSE7460_CTRL_VS_TGFB_TREATED_ACT_TCONV_DN, GSE7460_CTRL_VS_TGFB_TREATED_ACT_TCONV_DN GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_DN, GSE8921_UNSTIM_0H_VS_TLR1_2_STIM_MONOCYTE_12H_DN KASLER_HDAC7_TARGETS_1_UP, KASLER_HDAC7_TARGETS_1_UP GSE37301 PRO BCELL VS RAG2 KO NK CELL DN, GSE37301 PRO BCELL VS RAG2 KO NK CELL DN GSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_DN, GSE39820_CTRL_VS_IL1B_IL6_CD4_TCELL_DN GSE17721_12H_VS_24H_GARDIQUIMOD_BMDC_UP, GSE17721_12H_VS_24H_GARDIQUIMOD_BMDC_UP KEGG_COLORECTAL_CANCER, KEGG_COLORECTAL_CANCER KEGG PANCREATIC CANCER, KEGG PANCREATIC CANCER ATGCAGT_MIR217, ATGCAGT_MIR217 BASSO_CD40_SIGNALING_DN, BASSO_CD40_SIGNALING_DN KEGG_ADHERENS_JUNCTION, KEGG_ADHERENS_JUNCTION PID_IL2_1PATHWAY, PID_IL2_1PATHWAY GO_CELL_SUBSTRATE_ADHESION, GO_CELL_SUBSTRATE_ADHESION GO_KIDNEY_EPITHELIUM_DEVELOPMENT, GO_KIDNEY_EPITHELIUM_DEVELOPMENT PID_GMCSF_PATHWAY, PID_GMCSF_PATHWAY GO_NEPHRON_DEVELOPMENT, GO_NEPHRON_DEVELOPMENT SCHAEFFER_PROSTATE_DEVELOPMENT_12HR_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_12HR_UP CHEBOTAEV_GR_TARGETS_UP, CHEBOTAEV_GR_TARGETS_UP GO_REGULATION_OF_DENDRITE_DEVELOPMENT, GO_REGULATION_OF_DENDRITE_DEVELOPMENT VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_UP, GSE39820_TGFBETA1_IL6_VS_TGFBETA1_IL6_IL23A_TREATED_CD4_TCELL_UP GO_SKELETAL_SYSTEM_MORPHOGENESIS, GO_SKELETAL_SYSTEM_MORPHOGENESIS GO SMOOTH MUSCLE CELL DIFFERENTIATION, GO SMOOTH MUSCLE CELL DIFFERENTIATION GO_RENAL_TUBULE_DEVELOPMENT, GO_RENAL_TUBULE_DEVELOPMENT PID_NCADHERIN_PATHWAY, PID_NCADHERIN_PATHWAY GO_REGULATION_OF_DENDRITE_MORPHOGENESIS, GO_REGULATION_OF_DENDRITE_MORPHOGENESIS GO PROTEIN POLYMERIZATION, GO PROTEIN POLYMERIZATION GO_SYNAPTIC_VESICLE_CYCLE, GO_SYNAPTIC_VESICLE_CYCLE HNF3_Q6, HNF3_Q6 GO_VESICLE_DOCKING, GO_VESICLE_DOCKING GO_PROTEIN_SECRETION, GO_PROTEIN_SECRETION GO_NEURONAL_POSTSYNAPTIC_DENSITY, GO_NEURONAL_POSTSYNAPTIC_DENSITY GO_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION, GO_REGULATION_OF_T_HELPER_CELL_DIFFERENTIATION GO INTEGRATOR COMPLEX, GO INTEGRATOR COMPLEX LABBE_TGFB1_TARGETS_DN, LABBE_TGFB1_TARGETS_DN GO_ENDODERM_FORMATION, GO_ENDODERM_FORMATION GO_RAC_GTPASE_BINDING, GO_RAC_GTPASE_BINDING GO RESPONSE TO CALCIUM ION, GO RESPONSE TO CALCIUM ION PID_S1P_S1P4_PATHWAY, PID_S1P_S1P4_PATHWAY GO_DENDRITIC_SPINE_ORGANIZATION, GO_DENDRITIC_SPINE_ORGANIZATION GO_NEURON_FATE_COMMITMENT, GO_NEURON_FATE_COMMITMENT GO_REGULATION_OF_TOLERANCE_INDUCTION, GO_REGULATION_OF_TOLERANCE_INDUCTION GO_LAYER_FORMATION_IN_CEREBRAL_CORTEX, GO_LAYER_FORMATION_IN_CEREBRAL_CORTEX STAEGE_EWING_FAMILY_TUMOR, STAEGE_EWING_FAMILY_TUMOR ICHIBA_GRAFT_VERSUS_HOST_DISEASE_35D_DN, ICHIBA_GRAFT_VERSUS_HOST_DISEASE_35D_DN GO_LIPASE_ACTIVATOR_ACTIVITY, GO_LIPASE_ACTIVATOR_ACTIVITY

GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_UP, GSE3691_IFN_PRODUCING_KILLER_DC_VS_CONVENTIONAL_DC_SPLEEN_UP GSE39820 CTRL VS_TGFBETA1_IL6_CD4_TCELL_DN, GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_DN GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_PROGENITOR_DN, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_MONOCYTE_PROGENITOR_DN, GSE15330_MEGAKARYOCYTE_ERYTHROID_VS_GRANULOCYTE_PROGENITOR_DN, GSE15330_MEGAKARYOCYTE_PROGENITOR_DN, GSE15330 GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_UP, GSE15330_HSC_VS_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_IKAROS_KO_UP GSE15330 LYMPHOID MULTIPOTENT VS GRANULOCYTE MONOCYTE PROGENITOR IKAROS KO DN, GSE15330 LYMPHOID MULTIPOTENT VS GRANULOCYTE MC GSE16451_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_EQUINE_ENC_VIRUS_UP, GSE16451_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_EQUINE_E GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_HET_TCONV_DN, GSE7460_CTRL_VS_TGFB_TREATED_ACT_FOXP3_HET_TCONV_DN GSE24972_WT_VS_IRF8_KO_MARGINAL_ZONE_SPLEEN_BCELL_UP, GSE24972_WT_VS_IRF8_KO_MARGINAL_ZONE_SPLEEN_BCELL_UP GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_DN, GSE39916_B_CELL_SPLEEN_VS_PLASMA_CELL_BONE_MARROW_DN GSE36392_TYPE_2_MYELOID_VS_NEUTROPHIL_IL25_TREATED_LUNG_UP, GSE36392_TYPE_2_MYELOID_VS_NEUTROPHIL_IL25_TREATED_LUNG_UP GO_ESTABLISHMENT_OF_LOCALIZATION_BY_MOVEMENT_ALONG_MICROTUBULE, GO_ESTABLISHMENT_OF_LOCALIZATION_BY_MOVEMENT_ALONG_MICROTUBI GO CYTOSKELETON DEPENDENT INTRACELLULAR TRANSPORT, GO CYTOSKELETON DEPENDENT INTRACELLULAR TRANSPORT GSE4748_CTRL_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP, GSE4748_CTRL_VS_LPS_AND_CYANOBACTERIUM_LPSLIKE_STIM_DC_3H_UP GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN, GSE19941_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_DN HERNANDEZ ABERRANT MITOSIS BY DOCETACEL 2NM UP, HERNANDEZ ABERRANT MITOSIS BY DOCETACEL 2NM UP GO_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA, GO_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA GSE16450_CTRL_VS_IFNA_6H_STIM_MATURE_NEURON_CELL_LINE_UP, GSE16450_CTRL_VS_IFNA_6H_STIM_MATURE_NEURON_CELL_LINE_UP GO_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE, GO_BRANCHING_MORPHOGENESIS_OF_AN_EPITHELIAL_TUBE GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_UP, GSE19941_UNSTIM_VS_LPS_STIM_IL10_KO_MACROPHAGE_UP GO_NEGATIVE_REGULATION_OF_GLIAL_CELL_PROLIFERATION, GO_NEGATIVE_REGULATION_OF_GLIAL_CELL_PROLIFERATION GO POSITIVE REGULATION OF ENDOTHELIAL CELL DIFFERENTIATION, GO POSITIVE REGULATION OF ENDOTHELIAL CELL DIFFERENTIATION GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP, GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS, GO_DEVELOPMENTAL_GROWTH_INVOLVED_IN_MORPHOGENESIS GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION, GO_POSITIVE_REGULATION_OF_EPITHELIAL_CELL_DIFFERENTIATION GSE37416_CTRL_VS_0H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_0H_F_TULARENSIS_LVS_NEUTROPHIL_UP GO_BRANCHING_INVOLVED_IN_URETERIC_BUD_MORPHOGENESIS, GO_BRANCHING_INVOLVED_IN_URETERIC_BUD_MORPHOGENESIS SANCHEZ_MDM2_TARGETS, SANCHEZ_MDM2_TARGETS GO_ENDOCHONDRAL_BONE_MORPHOGENESIS, GO_ENDOCHONDRAL_BONE_MORPHOGENESIS