

G_DAY2_DERIVED_MACROPHAGE_DN, GSE1740_MCSF_VS_MCSF_AND_IFNG_DAY2_DERIVED_MACROPHAGE_DN

GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_T
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_12H_UP
GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_DN, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_T
AAGWWRNYGGC_UNKNOWN, AAGWWRNYGGC_UNKNOWN
GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_DN, GSE37301_MULTIPOTENT_PROGENITOR_VS_PRO_BCELL_DN
GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG
GSE13522_WT_VS_IFNAR_KO_SKIN_UP, GSE13522_WT_VS_IFNAR_KO_SKIN_UP
GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_UP, GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_UP
GO_PEPTIDYL_THREONINE_MODIFICATION, GO_PEPTIDYL_THREONINE_MODIFICATION
GSE22886_NAIVE_VS_IGG_IGA_MEMORY_BCELL_UP, GSE22886_NAIVE_VS_IGG_IGA_MEMORY_BCELL_UP
GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELL_UP
GO_WNT_SIGNALING_PATHWAY_CALCIUM_MODULATING_PATHWAY, GO_WNT_SIGNALING_PATHWAY_CALCIUM_MODU
GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP, GSE1791_CTRL_VS_NEUROMEDINU_IN_T_CELL_LINE_0.8H_UP
GO_RESPONSE_TO_MONOAMINE, GO_RESPONSE_TO_MONOAMINE
BECKER_TAMOXIFEN_RESISTANCE_DN, BECKER_TAMOXIFEN_RESISTANCE_DN
GO_REGULATION_OF_PROTEIN_MATURATION, GO_REGULATION_OF_PROTEIN_MATURATION
JEON_SMAD6_TARGETS_UP, JEON_SMAD6_TARGETS_UP
LIM_MAMMARY_LUMINAL_MATURE_UP, LIM_MAMMARY_LUMINAL_MATURE_UP
DOANE_BREAST_CANCER_CLASSES_UP, DOANE_BREAST_CANCER_CLASSES_UP
VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN, VANHARANTA_UTERINE_FIBROID_WITH_7Q_DELETION_DN
PID_ERBB4_PATHWAY, PID_ERBB4_PATHWAY
GO_REGULATION_OF_MICROTUBULE_POLYMERIZATION, GO_REGULATION_OF_MICROTUBULE_POLYMERIZATION
COATES_MACROPHAGE_M1_VS_M2_DN, COATES_MACROPHAGE_M1_VS_M2_DN
GO_PURINE_RIBONUCLEOSIDE_BISPHOSPHATE_METABOLIC_PROCESS, GO_PURINE_RIBONUCLEOSIDE_BISPHOSPHATE_MET
REACTOME_CYTOSOLIC_SULFONATION_OF_SMALL_MOLECULES, REACTOME_CYTOSOLIC_SULFONATION_OF_SMALL_MOL
GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_OT1_CD8AB_VS_HY_CD8AA_THYMOCYTE_
PID_SYNDECAN_1_PATHWAY, PID_SYNDECAN_1_PATHWAY
PAX6_01, PAX6_01
GO_NEGATIVE_REGULATION_OF_PROTEIN_MATURATION, GO_NEGATIVE_REGULATION_OF_PROTEIN_MATURATION
GO_REGULATION_OF_FATTY_ACID_TRANSPORT, GO_REGULATION_OF_FATTY_ACID_TRANSPORT
KEGG_SULFUR_METABOLISM, KEGG_SULFUR_METABOLISM
GO_SPHINGOLIPID_BIOSYNTHETIC_PROCESS, GO_SPHINGOLIPID_BIOSYNTHETIC_PROCESS
REACTOME_BIOLOGICAL_OXIDATIONS, REACTOME_BIOLOGICAL_OXIDATIONS
GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_VS_TGFB_AND_IL12_TREATED_ACT_CD4_
GO_CHONDROITIN_SULFATE_PROTEOGLYCAN_METABOLIC_PROCESS, GO_CHONDROITIN_SULFATE_PROTEOGLYCAN_MET