

1_VS_CPG_STIM_PDC_4H_UP, GSE7831_UNSTIM_VS_CPG_STIM_PDC_4H_UP

GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_MIR155_KO_UP, GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_MIR155_KO_UP
GSE6674_CPG_VS_CPG_AND_ANTI_IGM_STIM_BCELL_DN, GSE6674_CPG_VS_CPG_AND_ANTI_IGM_STIM_BCELL_DN
GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_UP, GSE11961_MARGINAL_ZONE_BCELL_VS_GERMINAL_CENTER_BCELL_DAY40_UP
KAECH_DAY8_EFF_VS_MEMORY_CD8_TCELL_DN, KAECH_DAY8_EFF_VS_MEMORY_CD8_TCELL_DN
GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_DN, GSE4590_LARGE_PRE_BCELL_VS_VPREB_POS_LARGE_PRE_BCELL_DN
GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_DN, GSE42021_CD24HI_TREG_VS_CD24HI_TCONV_THYMUS_DN
GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_UP, GSE43955_TH0_VS_TGFB_IL6_TH17_ACT_CD4_TCELL_4H_UP
GSE17721_LPS_VS_CPG_6H_BMDC_UP, GSE17721_LPS_VS_CPG_6H_BMDC_UP
GSE8384_CTRL_VS_B_ABORTUS_4H_MAC_CELL_LINE_DN, GSE8384_CTRL_VS_B_ABORTUS_4H_MAC_CELL_LINE_DN
UEDA_PERIFERAL_CLOCK, UEDA_PERIFERAL_CLOCK
LAKE_ADULT_KIDNEY_C6_PROXIMAL_TUBULE_EPITHELIAL_CELLS_FIBRINOGEN_POS_S3, LAKE_ADULT_KIDNEY_C6_PROXIMAL_TUBULE_EPITHELIAL_CELLS_FIBRINOGEN_POS_S3
LANDIS_ERBB2_BREAST_TUMORS_324_UP, LANDIS_ERBB2_BREAST_TUMORS_324_UP
WOOD_EBV_EBNA1_TARGETS_UP, WOOD_EBV_EBNA1_TARGETS_UP
GSE6259_DEC205_POS_DC_VS_CD8_TCELL_DN, GSE6259_DEC205_POS_DC_VS_CD8_TCELL_DN
JIANG_VHL_TARGETS, JIANG_VHL_TARGETS
GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_LOW_MONOCYTE_DN, GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_LOW_MONOCYTE_DN
GSE19888_ADENOSINE_A3R_INH_VS_INH_PRETREAT_AND_ACT_WITH_TCELL_MEMBRANES_MAST_CELL_DN, GSE19888_ADENOSINE_A3R_INH_VS_INH_PRETREAT_AND_ACT_WITH_TCELL_MEMBRANES_MAST_CELL_DN
GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_HEB_KO_DP_THYMOCYTE_UP
GOBP_MIRNA_METABOLIC_PROCESS, GOBP_MIRNA_METABOLIC_PROCESS
CHIANG_LIVER_CANCER_SUBCLASS_UNANNOTATED_UP, CHIANG_LIVER_CANCER_SUBCLASS_UNANNOTATED_UP
AGGGCCA_MIR328, AGGGCCA_MIR328
GOBP_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GOBP_REGULATION_OF_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
BIOCARTA_RANBP2_PATHWAY, BIOCARTA_RANBP2_PATHWAY
REACTOME_INTERACTION_BETWEEN_L1_AND_ANKYRINS, REACTOME_INTERACTION_BETWEEN_L1_AND_ANKYRINS
MODULE_213, MODULE_213
GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_MORPHOGENESIS, GOBP_APOPTOTIC_PROCESS_INVOLVED_IN_MORPHOGENESIS
GOBP_VASCULAR_WOUND_HEALING, GOBP_VASCULAR_WOUND_HEALING
GOBP_REGULATION_OF_MIRNA_METABOLIC_PROCESS, GOBP_REGULATION_OF_MIRNA_METABOLIC_PROCESS
BUSSLINGER_GASTRIC_NECK_CELLS, BUSSLINGER_GASTRIC_NECK_CELLS
MEBARKI_HCC_PROGENITOR_WNT_UP_CTNNB1_DEPENDENT, MEBARKI_HCC_PROGENITOR_WNT_UP_CTNNB1_DEPENDENT
GOBP_POTASSIUM_ION_HOMEOSTASIS, GOBP_POTASSIUM_ION_HOMEOSTASIS
GOBP_SODIUM_ION_HOMEOSTASIS, GOBP_SODIUM_ION_HOMEOSTASIS
REACTOME_DIGESTION_OF_DIETARY_LIPID, REACTOME_DIGESTION_OF_DIETARY_LIPID
HP_ABNORMAL_ST_SEGMENT, HP_ABNORMAL_ST_SEGMENT
GOCC_AXOLEMMA, GOCC_AXOLEMMA
BRIDEAU_IMPRINTED_GENES, BRIDEAU_IMPRINTED_GENES
MIR4462, MIR4462
GOBP_MONOVALENT_INORGANIC_ANION_HOMEOSTASIS, GOBP_MONOVALENT_INORGANIC_ANION_HOMEOSTASIS
GOMF_RNA_STEM_LOOP_BINDING, GOMF_RNA_STEM_LOOP_BINDING