

N_LPS_AND_IL10_STIM_MACROPHAGE_UP, GSE5589_UNSTIM_VS_180MIN_LPS_AND_IL10_STIM_MACROPHAGE_UP

- GSE5589_UNSTIM_VS_45MIN_LPS_AND_IL6_STIM_MACROPHAGE_DN, GSE5589_UNSTIM_VS_45MIN_LPS_AND_IL6_STIM_MACROPHAGE_DN
- GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN
- GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_STIM_MACROPHAGE_45MIN_DN
- GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN, GSE5589_WT_VS_IL6_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN
- GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN, GSE5589_IL6_KO_VS_IL10_KO_LPS_AND_IL10_STIM_MACROPHAGE_45MIN_DN
- GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN, GSE7460_WT_VS_FOXP3_HET_ACT_TCONV_DN
- GSE26890_CXCR1_NEG_VS_POS_EFFECTOR_CD8_TCELL_DN, GSE26890_CXCR1_NEG_VS_POS_EFFECTOR_CD8_TCELL_DN
- WAMUNYOKOLI_OVARIAN_CANCER_LMP_DN, WAMUNYOKOLI_OVARIAN_CANCER_LMP_DN
- GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_UP, GSE16522_ANTI_CD3CD28_STIM_VS_UNSTIM_MEMORY_CD8_TCELL_UP
- GSE5589_UNSTIM_VS_180MIN_LPS_STIM_MACROPHAGE_UP, GSE5589_UNSTIM_VS_180MIN_LPS_STIM_MACROPHAGE_UP
- MODULE_182, MODULE_182
- GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_DN, GSE13306_RA_VS_UNTREATED_MEM_CD4_TCELL_DN
- GSE30962_PRIMARY_VS_SECONDARY_ACUTE_LCMV_INF_CD8_TCELL_DN, GSE30962_PRIMARY_VS_SECONDARY_ACUTE_LCMV_INF_CD8_TCELL_DN
- GSE2405_0H_VS_3H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_UP, GSE2405_0H_VS_3H_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_UP
- KIM_MYCN_AMPLIFICATION_TARGETS_DN, KIM_MYCN_AMPLIFICATION_TARGETS_DN
- GSE4142_NAIVE_VS_GC_BCELL_UP, GSE4142_NAIVE_VS_GC_BCELL_UP
- GSE6875_TCONV_VS_TREG_UP, GSE6875_TCONV_VS_TREG_UP
- REACTOME_CELL_CELL_JUNCTION_ORGANIZATION, REACTOME_CELL_CELL_JUNCTION_ORGANIZATION
- TCCGTCC_MIR184, TCCGTCC_MIR184
- GO_CYCLOOXYGENASE_PATHWAY, GO_CYCLOOXYGENASE_PATHWAY
- GO_BONE_MINERALIZATION, GO_BONE_MINERALIZATION
- MANTOVANI_VIRAL_GPCR_SIGNALING_UP, MANTOVANI_VIRAL_GPCR_SIGNALING_UP
- GO_PHOSPHATIDYLETHANOLAMINE_BIOSYNTHETIC_PROCESS, GO_PHOSPHATIDYLETHANOLAMINE_BIOSYNTHETIC_PROCESS
- GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP, GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP
- FIGUEROA_AML_METHYLATION_CLUSTER_4_DN, FIGUEROA_AML_METHYLATION_CLUSTER_4_DN
- REACTOME_TIGHT_JUNCTION_INTERACTIONS, REACTOME_TIGHT_JUNCTION_INTERACTIONS
- GO_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS, GO_LONG_CHAIN_FATTY_ACID_METABOLIC_PROCESS
- GO_O_ACYLTRANSFERASE_ACTIVITY, GO_O_ACYLTRANSFERASE_ACTIVITY
- KEGG_ASCORBATE_AND_ALDARATE_METABOLISM, KEGG_ASCORBATE_AND_ALDARATE_METABOLISM
- CRX_DN.V1_DN, CRX_DN.V1_DN
- LUND_SILENCED_BY_METHYLATION, LUND_SILENCED_BY_METHYLATION
- HUANG_FOXA2_TARGETS_DN, HUANG_FOXA2_TARGETS_DN
- GO_ESTABLISHMENT_OF_NUCLEUS_LOCALIZATION, GO_ESTABLISHMENT_OF_NUCLEUS_LOCALIZATION