

NTI_IGM_STIM_TAK1_KO_BCELL_1H_UP, GSE41176_UNSTIM_VS_ANTI_IGM_STIM_TAK1_KO_BCELL_1H_UP

- GSE22919_RESTING_VS_IL2_IL12_IL15_STIM_NK_CELL_DN, GSE22919_RESTING_VS_IL2_IL12_IL15_STIM_NK_CELL_DN
- GSE27786_CD4_TCELL_VS_NKTCELL_UP, GSE27786_CD4_TCELL_VS_NKTCELL_UP
- GSE27786_CD8_TCELL_VS_NKTCELL_DN, GSE27786_CD8_TCELL_VS_NKTCELL_DN
- GSE37534_UNTREATED_VS_PIOGLITAZONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3_TRASDUCED_DN, GSE37534_UNTREATED_VS_PIOGLITAZONE_TREATED_CD4_TCELL_PPARG1_AND_FOXP3_TRASDUCED_DN
- GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP, GSE25123_IL4_VS_IL4_AND_ROSIGLITAZONE_STIM_MACROPHAGE_DAY10_UP
- GSE5455_HEALTHY_VS_TUMOR_BEARING_MOUSE_SPLEEN_MONOCYTE_24H_INCUBATION_DN, GSE5455_HEALTHY_VS_TUMOR_BEARING_MOUSE_SPLEEN_MONOCYTE_24H_INCUBATION_DN
- GSE25088_CTRL_VS_IL4_STIM_MACROPHAGE_DN, GSE25088_CTRL_VS_IL4_STIM_MACROPHAGE_DN
- GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_UP, GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_UP
- GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_UP, GSE17186_NAIVE_VS_CD21HIGH_TRANSITIONAL_BCELL_UP
- MODULE_171, MODULE_171
- HP_IMPAIRED_VIBRATORY_SENSATION, HP_IMPAIRED_VIBRATORY_SENSATION
- KESHELAVA_MULTIPLE_DRUG_RESISTANCE, KESHELAVA_MULTIPLE_DRUG_RESISTANCE
- GAVIN_FOXP3_TARGETS_CLUSTER_P7, GAVIN_FOXP3_TARGETS_CLUSTER_P7
- MIR4448, MIR4448
- LEE_LIVER_CANCER_E2F1_UP, LEE_LIVER_CANCER_E2F1_UP
- ZNF239_TARGET_GENES, ZNF239_TARGET_GENES
- CUI_DEVELOPING_HEART_LEFT_VENTRICULAR_CARDIOMYOCYTE, CUI_DEVELOPING_HEART_LEFT_VENTRICULAR_CARDIOMYOCYTE
- GOBP_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_LEUKOCYTE_APOPTOTIC_PROCESS
- HP_MACROGYRIA, HP_MACROGYRIA
- HP_ABNORMAL_ALDOLASE_LEVEL, HP_ABNORMAL_ALDOLASE_LEVEL
- GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_GLYCOPROTEIN, GOMF_CATALYTIC_ACTIVITY_ACTING_ON_A_GLYCOPROTEIN
- MIR3146, MIR3146
- HP_CEREBELLAR_DYSPLASIA, HP_CEREBELLAR_DYSPLASIA
- SINGH_KRAS_DEPENDENCY_SIGNATURE, SINGH_KRAS_DEPENDENCY_SIGNATURE
- GOBP_REGULATION_OF_LYMPHOCYTE_APOPTOTIC_PROCESS, GOBP_REGULATION_OF_LYMPHOCYTE_APOPTOTIC_PROCESS
- DALESSIO_TSA_RESPONSE, DALESSIO_TSA_RESPONSE
- HP_INTERRUPTED_AORTIC_ARCH, HP_INTERRUPTED_AORTIC_ARCH
- MODULE_543, MODULE_543
- GOCC_DENSE_CORE_GRANULE, GOCC_DENSE_CORE_GRANULE
- GOCC_AXOLEMMA, GOCC_AXOLEMMA
- GOBP_POSITIVE_REGULATION_OF_PROTEIN_MATURATION, GOBP_POSITIVE_REGULATION_OF_PROTEIN_MATURATION
- GOMF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS, GOMF_CYSSTEINE_TYPE_ENDOPEPTIDASE_ACTIVITY_INVOLVED_IN_APOPTOTIC_PROCESS
- MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_VS_CANDIN_PLACEBO_BCG_PRIMED_1DY_UP, MATSUMIYA_BLOOD_MODIFIED_VACCINIA_ANKARA_VACCINE_AGE_4_6MO_VACCINATED_VS_C
- HP_ILEUS, HP_ILEUS
- MUELLER_METHYLATED_IN_GLIOBLASTOMA, MUELLER_METHYLATED_IN_GLIOBLASTOMA
- GOBP_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS, GOBP_REGULATION_OF_T_CELL_DIFFERENTIATION_IN_THYMUS
- GOBP_POSITIVE_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS, GOBP_POSITIVE_REGULATION_OF_T_CELL_APOPTOTIC_PROCESS
- KRAS.50_UP.V1_DN, KRAS.50_UP.V1_DN
- GOBP_AMPA_GLUTAMATE_RECEPTOR_CLUSTERING, GOBP_AMPA_GLUTAMATE_RECEPTOR_CLUSTERING
- GOBP_REGULATION_OF_ENDOSOME_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT, GOBP_REGULATION_OF_ENDOSOME_TO_PLASMA_MEMBRANE_PROTEIN_TRANSPORT
- GOBP_TOLERANCE_INDUCTION_DEPENDENT_UPON_IMMUNE_RESPONSE, GOBP_TOLERANCE_INDUCTION_DEPENDENT_UPON_IMMUNE_RESPONSE
- GOBP_CELLULAR_RESPONSE_TO_AMMONIUM_ION, GOBP_CELLULAR_RESPONSE_TO_AMMONIUM_ION
- GOBP_REGULATION_OF_PRESYNAPTIC_MEMBRANE_POTENTIAL, GOBP_REGULATION_OF_PRESYNAPTIC_MEMBRANE_POTENTIAL
- GOBP_REGULATION_OF_GLUTAMATE_RECEPTOR_CLUSTERING, GOBP_REGULATION_OF_GLUTAMATE_RECEPTOR_CLUSTERING