

NTI_TREM1_AND_LPS_MONOCYTE_UP, GSE9988_ANTI_TREM1_VS_ANTI_TREM1_AND_LPS_MONOCYTE_UP

GSE16266_CTRL_VS_LPS_STIM_MEF_UP, GSE16266_CTRL_VS_LPS_STIM_MEF_UP
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE31082_DN_VS_CD4_SP_THYMOCYTE_DN, GSE31082_DN_VS_CD4_SP_THYMOCYTE_DN
GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP
GSE43863_TH1_VS_TFH_MEMORY_CD4_TCELL_DN, GSE43863_TH1_VS_TFH_MEMORY_CD4_TCELL_DN
GSE17721_LPS_VS_CPG_12H_BMDC_UP, GSE17721_LPS_VS_CPG_12H_BMDC_UP
GSE27434_WT_VS_DNMT1_KO_TREG_UP, GSE27434_WT_VS_DNMT1_KO_TREG_UP
GSE360_CTRL_VS_L_MAJOR_DC_UP, GSE360_CTRL_VS_L_MAJOR_DC_UP
GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_2H_DN, GSE36888_UNTREATED_VS_IL2_TREATED_TCELL_2H_DN
GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_DN
GSE30962_ACUTE_VS_CHRONIC_LCMV_SECONDARY_INF_CD8_TCELL_UP, GSE30962_ACUTE_VS_CHRONIC_LCMV_SECONDARY_INF_CD8_TCELL_UP
GSE3982_EFF_MEMORY_CD4_TCELL_VS_NKCELL_DN, GSE3982_EFF_MEMORY_CD4_TCELL_VS_NKCELL_DN
GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_CTRL_VS_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GO_RECEPTOR_SIGNALING_PROTEIN_ACTIVITY, GO_RECEPTOR_SIGNALING_PROTEIN_ACTIVITY
GSE17721_LPS_VS_POLYIC_12H_BMDC_DN, GSE17721_LPS_VS_POLYIC_12H_BMDC_DN
GSE43700_UNTREATED_VS_IL10_TREATED_PBMCDN, GSE43700_UNTREATED_VS_IL10_TREATED_PBMCDN
GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN, GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN
GNF2_HLA-C, GNF2_HLA-C
GSE43955_10H_VS_60H_ACT_CD4_TCELL_DN, GSE43955_10H_VS_60H_ACT_CD4_TCELL_DN
GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_UP, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_6H_UP
GSE21033_CTRL_VS_POLYIC_STIM_DC_24H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_24H_DN
GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_DN, GSE37605_C57BL6_VS_NOD_FOXP3_IRES_GFP_TCONV_DN
GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_UP
GO_REGULATION_OF_VACUOLAR_TRANSPORT, GO_REGULATION_OF_VACUOLAR_TRANSPORT
GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP, GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP
GSE7831_CPG_VS_INFLUENZA_STIM_PDC_4H_DN, GSE7831_CPG_VS_INFLUENZA_STIM_PDC_4H_DN
GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_DC_DN, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_DC_DN
GSE40441_NRP1_POS_INDUCED_TREG_VS_NRP1_NEG_NATURAL_TREG_DN, GSE40441_NRP1_POS_INDUCED_TREG_VS_NRP1_NEG_NATURAL_TREG_DN
GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
HOLLEMAN_DAUNORUBICIN_B_ALL_DN, HOLLEMAN_DAUNORUBICIN_B_ALL_DN
GO_TRANSLATION_REGULATOR_ACTIVITY, GO_TRANSLATION_REGULATOR_ACTIVITY
GO_MICROTUBULE_POLYMERIZATION, GO_MICROTUBULE_POLYMERIZATION
GSE14413_UNSTIM_VS_IFNB_STIM_L929_CELLS_UP, GSE14413_UNSTIM_VS_IFNB_STIM_L929_CELLS_UP
GO_CYTOKINE_BINDING, GO_CYTOKINE_BINDING
GO_MICROTUBULE_NUCLEATION, GO_MICROTUBULE_NUCLEATION
GO_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GO_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN, SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN
HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_DN, HOFMANN_MYELODYSPLASTIC_SYNDROM_RISK_DN
GO_NEGATIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_NEGATIVE_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
AZARE_STAT3_TARGETS, AZARE_STAT3_TARGETS
GO_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION, GO_POSITIVE_REGULATION_OF_INTERLEUKIN_4_PRODUCTION
PLASARI_TGFB1_TARGETS_1HR_DN, PLASARI_TGFB1_TARGETS_1HR_DN
ALONSO_METASTASIS_DN, ALONSO_METASTASIS_DN
PID_IL4_2PATHWAY, PID_IL4_2PATHWAY
GSE12198_LOW_IL2_STIM_NK_CELL_VS_HIGH_IL2_STIM_NK_CELL_DN, GSE12198_LOW_IL2_STIM_NK_CELL_VS_HIGH_IL2_STIM_NK_CELL_DN
GO_HISTONE_H2A_MONOUBIQUITINATION, GO_HISTONE_H2A_MONOUBIQUITINATION
ROYLANCE_BREAST_CANCER_16Q_COPY_NUMBER_UP, ROYLANCE_BREAST_CANCER_16Q_COPY_NUMBER_UP
GO_BLOC_COMPLEX, GO_BLOC_COMPLEX
GO_TRANSLATION_REPRESSOR_ACTIVITY, GO_TRANSLATION_REPRESSOR_ACTIVITY
GO_T_CELL_MIGRATION, GO_T_CELL_MIGRATION