

NTI_TREM1_AND_LPS_MONOCYTE_UP, GSE9988_ANTI_TREM1_VS_ANTI_TREM1_AND_LPS_MONOCYTE_UP

GSE2770_IL12_AND_TGFB_ACT_VS_GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN, GSE27859_MACROPHAGE_VS_CD11C_INT_F480_HI_MACROPHAGE_DN
GSE9988_ANTI_TREM1_VS_LOW_LPS_MONOCYTE_UP, GSE9988_ANTI_TREM1_VS_LOW_LPS_MONOCYTE_UP
GSE14769_UNSTIM_VS_60MIN_LPS_BMDM_UP, GSE14769_UNSTIM_VS_60MIN_LPS_BMDM_UP
GSE43863_TH1_VS_TFH_MEMORY_CD4_TCELL_DN, GSE43863_TH1_VS_TFH_MEMORY_CD4_TCELL_DN
GSE9988_LPS_VS_LPS_AND_ANTI_TREM1_MONOCYTE_DN, GSE9988_LPS_VS_LPS_AND_ANTI_TREM1_MONOCYTE_DN
GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PBMIC_UP, GSE9960_HEALTHY_VS_GRAM_POS_SEPSIS_PBMIC_UP
GSE15330_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_VS_PRO_BCELL_UP, GSE15330_MEGAKARYOCYTE_ERYTHROID_PROGENITOR_VS_PRO_BCELL_UP
GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_IFNG_PAM3CYS_IN_MAC_DN
GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_DN, GSE16385_ROSIGLITAZONE_VS_UNTREATED_IFNG_TNF_STIM_MACROPHAGE_DN
GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_DN
GSE37301_HEMATOPOIETIC_STEM_CELL_VS_LYMPHOID_PRIMED_MPP_UP, GSE37301_HEMATOPOIETIC_STEM_CELL_VS_LYMPHOID_PRIMED_MPP_UP
GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_ACT_VS_ACT_CD4_TCELL_2H_UP
GSE27434_WT_VS_DNMT1_KO_TREG_UP, GSE27434_WT_VS_DNMT1_KO_TREG_UP
GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_EOS_VS_FOXP3_AND_EOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
GSE17721_LPS_VS_POLYIC_12H_BMDC_DN, GSE17721_LPS_VS_POLYIC_12H_BMDC_DN
GNF2_HLA-C, GNF2_HLA-C
GSE18804_SPLEEN_MACROPHAGE_VS_COLON_TUMORAL_MACROPHAGE_UP, GSE18804_SPLEEN_MACROPHAGE_VS_COLON_TUMORAL_MACROPHAGE_UP
GSE36009_UNSTIM_VS_LPS_STIM_DC_UP, GSE36009_UNSTIM_VS_LPS_STIM_DC_UP
GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN, GSE17721_CTRL_VS_GARDIQUIMOD_0.5H_BMDC_DN
GNF2_SELL, GNF2_SELL
GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP, GSE1112_HY_CD8AB_VS_HY_CD8AA_THYMOCYTE_RTOC_CULTURE_UP
GSE21033_CTRL_VS_POLYIC_STIM_DC_24H_DN, GSE21033_CTRL_VS_POLYIC_STIM_DC_24H_DN
GO_ESTABLISHMENT_OF_LOCALIZATION_BY_MOVEMENT_ALONG_MICROTUBULE, GO_ESTABLISHMENT_OF_LOCALIZATION_BY_MOVEMENT_ALONG_MICROTUBULE
GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_DN, GSE22432_MULTIPOTENT_VS_COMMON_DC_PROGENITOR_DN
ACACTCC_MIR122A, ACACTCC_MIR122A
GO_REGULATION_OF_MEMBRANE_PERMEABILITY, GO_REGULATION_OF_MEMBRANE_PERMEABILITY
GNF2_MSN, GNF2_MSN
GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
PID_IL8_CXCR2_PATHWAY, PID_IL8_CXCR2_PATHWAY
GSE43700_UNTREATED_VS_IL10_TREATED_PBMIC_DN, GSE43700_UNTREATED_VS_IL10_TREATED_PBMIC_DN
GO_RECEPTOR_SIGNALING_PROTEIN_ACTIVITY, GO_RECEPTOR_SIGNALING_PROTEIN_ACTIVITY
GO_RAS_PROTEIN_SIGNAL_TRANSDUCTION, GO_RAS_PROTEIN_SIGNAL_TRANSDUCTION
chr1p34, chr1p34
GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_DC_DN, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_DC_DN
ATCTTGC_MIR31, ATCTTGC_MIR31
GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN, GSE339_CD4POS_VS_CD4CD8DN_DC_IN_CULTURE_DN
GO_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE, GO_ORGANELLE_TRANSPORT_ALONG_MICROTUBULE
MIF1_01, MIF1_01
GO_CELLULAR_PIGMENTATION, GO_CELLULAR_PIGMENTATION
CTCTGGA_MIR520A_MIR525, CTCTGGA_MIR520A_MIR525
GSE27786_BCELL_VS_NEUTROPHIL_DN, GSE27786_BCELL_VS_NEUTROPHIL_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
GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN, GSE33162_UNTREATED_VS_4H_LPS_STIM_HDAC3_HET_MACROPHAGE_DN
GO_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GO_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
RASHI_RESPONSE_TO_IONIZING_RADIATION_6, RASHI_RESPONSE_TO_IONIZING_RADIATION_6
GO_VESICLE_CYTOSKELETAL_TRAFFICKING, GO_VESICLE_CYTOSKELETAL_TRAFFICKING
GO_PIGMENTATION, GO_PIGMENTATION
GO_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX, GO_PHOSPHATIDYLINOSITOL_3_KINASE_COMPLEX
KESHELAVA_MULTIPLE_DRUG_RESISTANCE, KESHELAVA_MULTIPLE_DRUG_RESISTANCE
PID_KIT_PATHWAY, PID_KIT_PATHWAY
GSE22527_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_UP, GSE22527_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_UP
NAKAMURA_METASTASIS, NAKAMURA_METASTASIS
GO_AXO_DENDRITIC_TRANSPORT, GO_AXO_DENDRITIC_TRANSPORT
GO_PLATELET_AGGREGATION, GO_PLATELET_AGGREGATION
GO_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION, GO_REGULATION_OF_LYMPHOCYTE_DIFFERENTIATION
GO_MICROTUBULE_NUCLEATION, GO_MICROTUBULE_NUCLEATION
GO_AXON_CYTOPLASM, GO_AXON_CYTOPLASM
GO_REGULATION_OF_VACUOLAR_TRANSPORT, GO_REGULATION_OF_VACUOLAR_TRANSPORT
GO_HISTONE_UBIQUITINATION, GO_HISTONE_UBIQUITINATION
GO_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY, GO_PROTEIN_SERINE_THREONINE_PHOSPHATASE_ACTIVITY
GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_UP, GSE11961_FOLLICULAR_BCELL_VS_MARGINAL_ZONE_BCELL_UP
CREIGHTON_AKT1_SIGNALING_VIA_MTOR_DN, CREIGHTON_AKT1_SIGNALING_VIA_MTOR_DN
VARELA_ZMPSTE24_TARGETS_UP, VARELA_ZMPSTE24_TARGETS_UP
BIOCARTA_GH_PATHWAY, BIOCARTA_GH_PATHWAY
GO_HOMOTYPIC_CELL_CELL_ADHESION, GO_HOMOTYPIC_CELL_CELL_ADHESION
GSE14413_UNSTIM_VS_IFNB_STIM_1929_CELLS_UP, GSE14413_UNSTIM_VS_IFNB_STIM_1929_CELLS_UP
GO_HISTONE_MONOUBIQUITINATION, GO_HISTONE_MONOUBIQUITINATION
GO_CLATHRIN_COAT, GO_CLATHRIN_COAT
GO_CELL_PROJECTION_CYTOPLASM, GO_CELL_PROJECTION_CYTOPLASM
GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_129_MOUSE_DN, GSE13522_CTRL_VS_T_CRUZI_Y_STRAIN_INF_SKIN_129_MOUSE_DN
GO_NEGATIVE_REGULATION_OF_INNATE_IMMUNE_RESPONSE, GO_NEGATIVE_REGULATION_OF_INNATE_IMMUNE_RESPONSE
GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GO_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS
GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_UP, GSE1740_UNSTIM_VS_IFNA_STIMULATED_MCSF_IFNG_DERIVED_MACROPHAGE_UP
GO_CARBOHYDRATE_BIOSYNTHETIC_PROCESS, GO_CARBOHYDRATE_BIOSYNTHETIC_PROCESS
HOLLEMAN_DAUNORUBICIN_ALL_DN, HOLLEMAN_DAUNORUBICIN_ALL_DN
GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_UP, GSE21670_UNTREATED_VS_TGFB_TREATED_STAT3_KO_CD4_TCELL_UP
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
SHIPP_DLBCL_VS_FOLLICULAR_LYMPHOMA_DN, SHIPP_DLBCL_VS_FOLLICULAR_LYMPHOMA_DN
GO_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION, GO_NEGATIVE_REGULATION_OF_TUMOR_NECROSIS_FACTOR_SUPERFAMILY_CYTOKINE_PRODUCTION
GO_CLATHRIN_ADAPTOR_COMPLEX, GO_CLATHRIN_ADAPTOR_COMPLEX
REACTOME_NUCLEAR_RECEPTOR_TRANSCRIPTION_PATHWAY, REACTOME_NUCLEAR_RECEPTOR_TRANSCRIPTION_PATHWAY
GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION, GO_NEGATIVE_REGULATION_OF_MITOCHONDRION_ORGANIZATION
GO_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_LIPID_BIOSYNTHETIC_PROCESS
chr15q22, chr15q22
GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP, GSE15659_NAIVE_CD4_TCELL_VS_NONSUPPRESSIVE_TCELL_UP
GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_LOW_MONOCYTE_DN, GSE32034_UNTREATED_VS_ROSIGLIZATONE_TREATED_LY6C_LOW_MONOCYTE_DN
GO_REGULATION_OF_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_PROTEIN_INSERTION_INTO_MITOCHONDRIAL_MEMBRANE_INVOLVED_IN_APOPTOTIC_SIGNALING_PATHWAY
GO_NEGATIVE_REGULATION_OF_NEURON_DIFFERENTIATION, GO_NEGATIVE_REGULATION_OF_NEURON_DIFFERENTIATION
GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_DN, GARGALOVIC_RESPONSE_TO_OXIDIZED_PHOSPHOLIPIDS_RED_DN
SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN, SARTIPY_NORMAL_AT_INSULIN_RESISTANCE_DN
GO_HISTONE_H2A_UBIQUITINATION, GO_HISTONE_H2A_UBIQUITINATION
REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS, REACTOME_APOPTOTIC_CLEAVAGE_OF_CELLULAR_PROTEINS
SIG_INSULIN_RECEPTOR_PATHWAY_IN_CARDIAC_MYOCYTES, SIG_INSULIN_RECEPTOR_PATHWAY_IN_CARDIAC_MYOCYTES
GO_LUMENAL_SIDE_OF_MEMBRANE, GO_LUMENAL_SIDE_OF_MEMBRANE
GO_ARF_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY, GO_ARF_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_ACTIVITY
HFH3_01, HFH3_01
LEE_CALORIE_RESTRICTION_MUSCLE_UP, LEE_CALORIE_RESTRICTION_MUSCLE_UP
GO_RACEMASE_AND_EPIMERASE_ACTIVITY, GO_RACEMASE_AND_EPIMERASE_ACTIVITY
GO_PIGMENT GRANULE_ORGANIZATION, GO_PIGMENT GRANULE_ORGANIZATION
GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY, GO_TRANSMEMBRANE_RECEPTOR_PROTEIN_SERINE_THREONINE_KINASE_SIGNALING_PATHWAY
GSE43956_WT_VS_SGK1_KO_THI17_DIFFERENTIATED_CD4_TCELL_UP, GSE43956_WT_VS_SGK1_KO_THI17_DIFFERENTIATED_CD4_TCELL_UP
GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP, GSE15659_NONSUPPRESSIVE_TCELL_VS_ACTIVATED_TREG_UP
PID_IL4_2PATHWAY, PID_IL4_2PATHWAY
GO_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT, GO_SYNAPTIC_VESICLE_CYTOSKELETAL_TRANSPORT
REACTOME_G_BETA_GAMMA_SIGNALLING_THROUGH_PI3KGAMMA, REACTOME_G_BETA_GAMMA_SIGNALLING_THROUGH_PI3KGAMMA
FONTAINE_FOLLICULAR_THYROID_ADENOMA_DN, FONTAINE_FOLLICULAR_THYROID_ADENOMA_DN