

CT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_24H_CD4_TCELL_DN

GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_4H_CD4_TCELL_DN
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_2H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_2H_CD4_TCELL_DN
GSE17974_0H_VS_48H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_48H_IN_VITRO_ACT_CD4_TCELL_DN
GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_12H_IN_VITRO_ACT_CD4_TCELL_DN
GSE16451_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_EQUINE_ENC_VIRUS_DN, GSE16451_IMMATURE_VS_MATURE_NEURON_CELL_LINE_WEST_EQUINE_ENC_VIRUS_DN
GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_UP, GSE22033_UNTREATED_VS_ROSIGLITAZONE_TREATED_MEF_UP
GSE22103_LPS_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP, GSE22103_LPS_VS_GMCSF_AND_IFNG_STIM_NEUTROPHIL_UP
GSE19888_ADENOSINE_A3R_ACT_VS_A3R_ACT_WITH_A3R_INH_PRETREATMENT_IN_MAST_CELL_UP, GSE19888_ADENOSINE_A3R_ACT_VS_A3R_ACT_WITH_A3R_INH_PRETREATMENT_IN_MAST_CELL_UP
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_72H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_72H_CD4_TCELL_DN
GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_2H_IN_VITRO_ACT_CD4_TCELL_DN
GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_DN, GSE17974_2H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_DN
GSE19923_WT_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_WT_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
GSE17974_1.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_DN, GSE17974_1.5H_VS_72H_IL4_AND_ANTI_IL12_ACT_CD4_TCELL_DN
GSE6674_CPG_VS_PL2_3_STIM_BCELL_UP, GSE6674_CPG_VS_PL2_3_STIM_BCELL_UP
GSE13738_RESTING_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN, GSE13738_RESTING_VS_BYSTANDER_ACTIVATED_CD4_TCELL_DN
GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_20MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_20MIN_DN
GSE27786_LSK_VS_NKCELL_UP, GSE27786_LSK_VS_NKCELL_UP
GSE17974_0H_VS_1H_IN_VITRO_ACT_CD4_TCELL_DN, GSE17974_0H_VS_1H_IN_VITRO_ACT_CD4_TCELL_DN
GSE3982_BASOPHIL_VS_TH1_DN, GSE3982_BASOPHIL_VS_TH1_DN
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_6H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_6H_CD4_TCELL_DN
GSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_DN, GSE8621_UNSTIM_VS_LPS_PRIMED_UNSTIM_MACROPHAGE_DN
GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP, GSE10273_HIGH_IL7_VS_HIGH_IL7_AND_IRF4_IN_IRF4_8_NULL_PRE_BCELL_UP
GSE14308_TH17_VS_NATURAL_TREG_UP, GSE14308_TH17_VS_NATURAL_TREG_UP
GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN, GSE17974_CTRL_VS_ACT_IL4_AND_ANTI_IL12_1H_CD4_TCELL_DN
PRC2_EED_UP.V1_DN, PRC2_EED_UP.V1_DN
GSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN, GSE11864_UNTREATED_VS_CSF1_PAM3CYS_IN_MAC_DN
GSE22033_WT_VS_PPARG_KO_MEF_DN, GSE22033_WT_VS_PPARG_KO_MEF_DN
GSE11924_TFH_VS_TH17_CD4_TCELL_DN, GSE11924_TFH_VS_TH17_CD4_TCELL_DN
GSE9650_NAIVE_VS_EFF_CD8_TCELL_DN, GSE9650_NAIVE_VS_EFF_CD8_TCELL_DN
CHUNG_BLISTER_CYTOTOXICITY_UP, CHUNG_BLISTER_CYTOTOXICITY_UP
GSE5679_CTRL_VS_PPARG_LIGAND_ROSIGLITAZONE_AND_RARA_AAGONIST_AM580_TREATED_DC_DN, GSE5679_CTRL_VS_PPARG_LIGAND_ROSIGLITAZONE_AND_RARA_AAGONIST_AM580_TREATED_DC_DN
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMIC_AT_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMIC_AT_DX_DN
COLDREN_GEFITINIB_RESISTANCE_UP, COLDREN_GEFITINIB_RESISTANCE_UP
GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_NAIVE_VS_DAY6_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_DN, GSE24142_EARLY_THYMIC_PROGENITOR_VS_DN2_THYMOCYTE_DN
GSE21927_SPLEEN_VS_4T1_TUMOR_MONOCYTE_BALBC_DN, GSE21927_SPLEEN_VS_4T1_TUMOR_MONOCYTE_BALBC_DN
GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN, GSE20152_HTNFA_OVERXPRESS_ANKLE_VS_CTRL_SPHK1_KO_ANKLE_DN
GSE5679_RARA_AAGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP, GSE5679_RARA_AAGONIST_AM580_VS_AM580_AND_ROSIGLITAZONE_TREATED_DC_UP
GSE4811_CLASSICALY_ACTIVATED_VS_TYPE_2_ACTIVATED_MACROPHAGE_UP, GSE4811_CLASSICALY_ACTIVATED_VS_TYPE_2_ACTIVATED_MACROPHAGE_UP
GSE17580_TREG_VS_TEFF_UP, GSE17580_TREG_VS_TEFF_UP
ESC_J1_UP_LATE.V1_DN, ESC_J1_UP_LATE.V1_DN
MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_DN, MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_DN
PASINI_SUZ12_TARGETS_UP, PASINI_SUZ12_TARGETS_UP
GSE15659_RESTING_TREG_VS_NONSUPPRESSIVE_TCELL_DN, GSE15659_RESTING_TREG_VS_NONSUPPRESSIVE_TCELL_DN
GO_RNA_POLYMERASE_ACTIVITY, GO_RNA_POLYMERASE_ACTIVITY
DARWICHE_PAPILLOMA_RISK_LOW_DN, DARWICHE_PAPILLOMA_RISK_LOW_DN
GSE8835_HEALTHY_VS_CLL_CD4_TCELL_UP, GSE8835_HEALTHY_VS_CLL_CD4_TCELL_UP
GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMIC_4MONTH_POST_DX_DN, GSE9006_HEALTHY_VS_TYPE_1_DIABETES_PBMIC_4MONTH_POST_DX_DN
GO_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_NITROGEN_BONDS
ZHAN_V2_LATE_DIFFERENTIATION_GENES, ZHAN_V2_LATE_DIFFERENTIATION_GENES
GO_POSITIVE_REGULATION_OF_HISTONE_METHYLATION, GO_POSITIVE_REGULATION_OF_HISTONE_METHYLATION
GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN, GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN
GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDONUCLEASE_ACTIVITY_ACTIVE_WITH_EITHER_RIBO_OR_DEOXYRIBONUCLEIC_ACIDS_AND_PRODUCING_5_PHOSPHOMONOESTERS
LIU_PROSTATE_CANCER_UP, LIU_PROSTATE_CANCER_UP
GSE17721_LPS_VS_CPG_24H_BMDC_UP, GSE17721_LPS_VS_CPG_24H_BMDC_UP
REACTOME_METABOLISM_OF_VITAMINS_AND_COFACTORS, REACTOME_METABOLISM_OF_VITAMINS_AND_COFACTORS
GSE26912_TUMORICIDAL_VS_CTRL_MACROPHAGE_DN, GSE26912_TUMORICIDAL_VS_CTRL_MACROPHAGE_DN
GO_MEDIATOR_COMPLEX, GO_MEDIATOR_COMPLEX
GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN, GSE36891_POLYIC_TLR3_VS_PAM_TLR2_STIM_PERITONEAL_MACROPHAGE_DN
GO_HOMEOSTASIS_OF_NUMBER_OF_CELLS, GO_HOMEOSTASIS_OF_NUMBER_OF_CELLS
CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP, CHEOK_RESPONSE_TO_MERCAPTOPURINE_UP
GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_DN, GSE17580_UNINFECTED_VS_S_MANSONI_INF_TEFF_DN
GO_EXODEOXYRIBONUCLEASE_ACTIVITY, GO_EXODEOXYRIBONUCLEASE_ACTIVITY
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_DN, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_DN
GO_ENDORIBONUCLEASE_ACTIVITY, GO_ENDORIBONUCLEASE_ACTIVITY
GO_CENTROSOME_LOCALIZATION, GO_CENTROSOME_LOCALIZATION
NIKOLSKY_BREAST_CANCER_11Q12_Q14_AMPLICON, NIKOLSKY_BREAST_CANCER_11Q12_Q14_AMPLICON
GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER, GO_REGULATION_OF_TRANSCRIPTION_ELONGATION_FROM_RNA_POLYMERASE_II_PROMOTER
BILBAN_B_CLL_LPL_UP, BILBAN_B_CLL_LPL_UP
GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_CTRL_VS_48H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GSE21927_SPLEEN_VS_TUMOR_MONOCYTE_BALBC_UP, GSE21927_SPLEEN_VS_TUMOR_MONOCYTE_BALBC_UP
REACTOME_SYNTHESIS_OF_PIP3_AT_THE_EARLY_ENDOSOME_MEMBRANE, REACTOME_SYNTHESIS_OF_PIP3_AT_THE_EARLY_ENDOSOME_MEMBRANE
GAUSSMANN_MLL_AF4_FUSION_TARGETS_E_DN, GAUSSMANN_MLL_AF4_FUSION_TARGETS_E_DN
AR_01, AR_01
GO_PTERIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_PTERIDINE_CONTAINING_COMPOUND_METABOLIC_PROCESS
GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDORIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
GO_FOLIC_ACID_CONTAINING_COMPOUND_METABOLIC_PROCESS, GO_FOLIC_ACID_CONTAINING_COMPOUND_METABOLIC_PROCESS
MURAKAMI_UV_RESPONSE_6HR_DN, MURAKAMI_UV_RESPONSE_6HR_DN
GO_POLYOL_CATABOLIC_PROCESS, GO_POLYOL_CATABOLIC_PROCESS
GO_CELLULAR_RESPONSE_TO_OXYGEN_LEVELS, GO_CELLULAR_RESPONSE_TO_OXYGEN_LEVELS
GTCNYYATGR_UNKNOWN, GTCNYYATGR_UNKNOWN
GAZIN_EPIGENETIC_SILENCING_BY_KRAS, GAZIN_EPIGENETIC_SILENCING_BY_KRAS
GO_DNA_HELICASE_COMPLEX, GO_DNA_HELICASE_COMPLEX
GO_NEURON_DEATH, GO_NEURON_DEATH
GO_REGULATION_OF_EXIT_FROM_MITOSIS, GO_REGULATION_OF_EXIT_FROM_MITOSIS
GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP
GO_INOSITOL_PHOSPHATE_CATABOLIC_PROCESS, GO_INOSITOL_PHOSPHATE_CATABOLIC_PROCESS
LOPEZ_MESOTELIOMA_SURVIVAL_TIME_UP, LOPEZ_MESOTELIOMA_SURVIVAL_TIME_UP
FIGUEROA_AML_METHYLATION_CLUSTER_2_UP, FIGUEROA_AML_METHYLATION_CLUSTER_2_UP
chr4p15, chr4p15
GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR, GO_OXIDOREDUCTASE_ACTIVITY_ACTING_ON_THE_CH_NH_GROUP_OF_DONORS_NAD_OR_NADP_AS_ACCEPTOR
GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_UP, GSE17721_CPG_VS_GARDIQUIMOD_16H_BMDC_UP
GO_NEURON_APOPTOTIC_PROCESS, GO_NEURON_APOPTOTIC_PROCESS
GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_DN, GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_DN
GO_FOLIC_ACID_METABOLIC_PROCESS, GO_FOLIC_ACID_METABOLIC_PROCESS
CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB, CARDOSO_RESPONSE_TO_GAMMA_RADIATION_AND_3AB
WEST_ADRENOCORTICAL_CARCINOMA_VS_ADENOMA_UP, WEST_ADRENOCORTICAL_CARCINOMA_VS_ADENOMA_UP