

ED_RNA_VS_MRNA_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_DC_UP

GSE14000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_4H_LPS_DC_UP
GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_UP, GSE14000_UNSTIM_VS_4H_LPS_DC_TRANSLATED_RNA_UP
GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_DN, GSE2770_IL12_AND_TGFB_ACT_VS_ACT_CD4_TCELL_48H_DN
GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_UP, GSE41867_NAIVE_VS_DAY30_LCMV_ARMSTRONG_MEMORY_CD8_TCELL_UP
GO_MACROMOLECULAR_COMPLEX_DISASSEMBLY, GO_MACROMOLECULAR_COMPLEX_DISASSEMBLY
GSE27241_WT_VS_RORGT_KO_TH17_POLARIZED_CD4_TCELL_DN, GSE27241_WT_VS_RORGT_KO_TH17_POLARIZED_CD4_TCELL_DN
GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP, GSE37301_LYMPHOID_PRIMED_MPP_VS_RAG2_KO_NK_CELL_UP
GSE3982_MAST_CELL_VS_EFF_MEMORY_CD4_TCELL_UP, GSE3982_MAST_CELL_VS_EFF_MEMORY_CD4_TCELL_UP
GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP, GSE2770_IL12_AND_TGFB_VS_IL4_TREATED_ACT_CD4_TCELL_2H_UP
GSE14000_UNSTIM_VS_16H_LPS_DC_DN, GSE14000_UNSTIM_VS_16H_LPS_DC_DN
GGAANCGGAANY_UNKNOWN, GGAANCGGAANY_UNKNOWN
GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_48H_UP
GSE6269_HEALTHY_VS_FLU_INF_PBM_C_UP, GSE6269_HEALTHY_VS_FLU_INF_PBM_C_UP
GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_UP, GSE39820_CTRL_VS_TGFBETA1_IL6_CD4_TCELL_UP
GO_CYTOPLASMIC_MRNA_PROCESSING_BODY, GO_CYTOPLASMIC_MRNA_PROCESSING_BODY
GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS, GO_LIGASE_ACTIVITY_FORMING_CARBON_OXYGEN_BONDS
GSE2405_HEAT_KILLED_LYSATE_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_9H_DN, GSE2405_HEAT_KILLED_LYSATE_VS_LIVE_A_PHAGOCYTOPHILUM_STIM_NEUTROPHIL_9H_DN
GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY, GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_ASSEMBLY
GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_DN, GSE10856_CTRL_VS_TNFRSF6B_IN_MACROPHAGE_DN
GSE25087_TREG_VS_TCONV_FETUS_DN, GSE25087_TREG_VS_TCONV_FETUS_DN
GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_DN
GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN, GSE21670_UNTREATED_VS_TGFB_IL6_TREATED_CD4_TCELL_DN
GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_I_BIOGENESIS, GO_MITOCHONDRIAL_RESPIRATORY_CHAIN_COMPLEX_I_BIOGENESIS
GSE45365_NK_CELL_VS_CD11B_DC_DN, GSE45365_NK_CELL_VS_CD11B_DC_DN
GO_PROTEIN_N_TERMINUS_BINDING, GO_PROTEIN_N_TERMINUS_BINDING
GO_SMALL_SUBUNIT_PROCESSOME, GO_SMALL_SUBUNIT_PROCESSOME
GO_SWI_SNF_SUPERFAMILY_TYPE_COMPLEX, GO_SWI_SNF_SUPERFAMILY_TYPE_COMPLEX
GO_TRANSLATION_INITIATION_FACTOR_BINDING, GO_TRANSLATION_INITIATION_FACTOR_BINDING
GO_RNA_POLYMERASE_BINDING, GO_RNA_POLYMERASE_BINDING
GSE37534_GW1929_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP, GSE37534_GW1929_VS_ROSIGLITAZONE_TREATED_CD4_TCELL_PPARG1_FOXP3_TRANSDUCED_UP
GO_INO80_TYPE_COMPLEX, GO_INO80_TYPE_COMPLEX
GO_EXOSOME_RNASE_COMPLEX, GO_EXOSOME_RNASE_COMPLEX
GO_RRNA_MODIFICATION, GO_RRNA_MODIFICATION
GO_RNA_POLYMERASE_CORE_ENZYME_BINDING, GO_RNA_POLYMERASE_CORE_ENZYME_BINDING
GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING, GO_BASAL_TRANSCRIPTION_MACHINERY_BINDING
AKT_UP_MTOR_DN.V1_DN, AKT_UP_MTOR_DN.V1_DN
GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN, GSE22443_NAIVE_VS_ACT_AND_IL12_TREATED_CD8_TCELL_DN
GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN, GSE21927_EL4_VS_MCA203_TUMOR_MONOCYTES_DN
GO_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX, GO_TRANSCRIPTION_ELONGATION_FACTOR_COMPLEX
GO_ORGANELLAR_SMALL_RIBOSOMAL_SUBUNIT, GO_ORGANELLAR_SMALL_RIBOSOMAL_SUBUNIT
GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_UP, GSE26488_WT_VS_HDAC7_DELTAP_TG_OT2_THYMOCYTE_WITH_PEPTIDE_INJECTION_UP
KORKOLA_EMBRYONAL_CARCINOMA_UP, KORKOLA_EMBRYONAL_CARCINOMA_UP
REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS, REACTOME_GLYCOGEN_BREAKDOWN_GLYCOGENOLYSIS
GO_DEMETHYLASE_ACTIVITY, GO_DEMETHYLASE_ACTIVITY
DAIRKEE_TERT_TARGETS_DN, DAIRKEE_TERT_TARGETS_DN
GO_DEMETHYLATION, GO_DEMETHYLATION
BURTON_ADIPOGENESIS_10, BURTON_ADIPOGENESIS_10
GO_VENTRICULAR_SYSTEM_DEVELOPMENT, GO_VENTRICULAR_SYSTEM_DEVELOPMENT
GO_HISTONE_MONOUBIQUITINATION, GO_HISTONE_MONOUBIQUITINATION
GO_DNA_HELICASE_COMPLEX, GO_DNA_HELICASE_COMPLEX
GO_RNA_SURVEILLANCE, GO_RNA_SURVEILLANCE
chr3q26, chr3q26
GO_RESPONSE_TO_HEAT, GO_RESPONSE_TO_HEAT
GO_TRANSCRIPTION_EXPORT_COMPLEX, GO_TRANSCRIPTION_EXPORT_COMPLEX
LI_CYTIDINE_ANALOGS_CYCOTOTOXICITY, LI_CYTIDINE_ANALOGS_CYCOTOTOXICITY
GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP, GSE14000_TRANSLATED_RNA_VS_MRNA_16H_LPS_DC_UP
GO_AMINOACYL_TRNA_EDITING_ACTIVITY, GO_AMINOACYL_TRNA_EDITING_ACTIVITY
PAX5_01, PAX5_01
GO_GLUCAN_METABOLIC_PROCESS, GO_GLUCAN_METABOLIC_PROCESS
GO_RRNA_METHYLATION, GO_RRNA_METHYLATION
GO_LATERAL_VENTRICLE_DEVELOPMENT, GO_LATERAL_VENTRICLE_DEVELOPMENT
GSE43957_UNTREATED_VS_NACL_TREATED_ANTI_CD3_CD28_STIM_CD4_TCELL_UP, GSE43957_UNTREATED_VS_NACL_TREATED_ANTI_CD3_CD28_STIM_CD4_TCELL_UP
GO_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT, GO_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_TRANSPORT
GO_PROTEASOME_BINDING, GO_PROTEASOME_BINDING
GO_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_TRANSLATION_IN_RESPONSE_TO_STRESS
GO_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION, GO_POSITIVE_REGULATION_OF_NEUROTRANSMITTER_SECRETION
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX6_DN, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX6_DN
GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY, GO_PHOSPHATIDYLINOSITOL_PHOSPHATE_PHOSPHATASE_ACTIVITY
GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_DN, GSE23505_UNTREATED_VS_4DAY_IL6_IL1_TREATED_CD4_TCELL_DN
GO_REGULATION_OF_TRANSLATIONAL_INITIATION_IN_RESPONSE_TO_STRESS, GO_REGULATION_OF_TRANSLATIONAL_INITIATION_IN_RESPONSE_TO_STRESS
GO_PHOSPHOLIPID_DEPHOSPHORYLATION, GO_PHOSPHOLIPID_DEPHOSPHORYLATION
GO_OLIGODENDROCYTE_DEVELOPMENT, GO_OLIGODENDROCYTE_DEVELOPMENT
GO_MITOCHONDRIAL_DNA_METABOLIC_PROCESS, GO_MITOCHONDRIAL_DNA_METABOLIC_PROCESS
GSE17721_PAM3CSK4_VS_CPG_2H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_2H_BMDC_DN
GO_GLYCOSIDE_METABOLIC_PROCESS, GO_GLYCOSIDE_METABOLIC_PROCESS
PPARG_01, PPARG_01
GO_SWI_SNF_COMPLEX, GO_SWI_SNF_COMPLEX
GO_GLUCAN_BIOSYNTHETIC_PROCESS, GO_GLUCAN_BIOSYNTHETIC_PROCESS
GO_TRANSCRIPTION_FROM_MITOCHONDRIAL_PROMOTER, GO_TRANSCRIPTION_FROM_MITOCHONDRIAL_PROMOTER
POS_HISTAMINE_RESPONSE_NETWORK, POS_HISTAMINE_RESPONSE_NETWORK