

_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN

GSE2770_TGFB_AND_IL4_VS_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN
GSE17974_1H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP, GSE17974_1H_VS_72H_UNTREATED_IN_VITRO_CD4_TCELL_UP
GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN
REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE, REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE
GSE42724_MEMORY_VS_B1_BCELL_UP, GSE42724_MEMORY_VS_B1_BCELL_UP
GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY10_DN, GSE24634_IL4_VS_CTRL_TREATED_NAIVE_CD4_TCELL_DAY10_DN
GSE14769_UNSTIM_VS_120MIN_LPS_BMDM_DN, GSE14769_UNSTIM_VS_120MIN_LPS_BMDM_DN
HALLMARK_UV_RESPONSE_UP, HALLMARK_UV_RESPONSE_UP
GO_POSITIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY, GO_POSITIVE_REGULATION_OF_APOPTOTIC_SIGNALING_PATHWAY
PHONG_TNF_RESPONSE_VIA_P38_PARTIAL, PHONG_TNF_RESPONSE_VIA_P38_PARTIAL
KYNG_DNA_DAMAGE_DN, KYNG_DNA_DAMAGE_DN
ELVIDGE_HYPOXIA_BY_DMOG_UP, ELVIDGE_HYPOXIA_BY_DMOG_UP
GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_UP, GSE12392_CD8A_POS_VS_NEG_SPLEEN_DC_UP
GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP, GSE43955_TH0_VS_TGFB_IL6_IL23_TH17_ACT_CD4_TCELL_60H_UP
GSE3982_CTRL_VS_LPS_48H_DC_DN, GSE3982_CTRL_VS_LPS_48H_DC_DN
GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_AND_P13K_INHIBITION_UP, GSE19772_CTRL_VS_HCMV_INF_MONOCYTES_AND_P13K_INHIBITION_UP
GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP, GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP
MENSE_HYPOXIA_UP, MENSE_HYPOXIA_UP
PEART_HDAC_PROLIFERATION_CLUSTER_UP, PEART_HDAC_PROLIFERATION_CLUSTER_UP
GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_UP, GSE360_L_MAJOR_VS_B_MALAYI_LOW_DOSE_DC_UP
GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_DN, GSE22033_UNTREATED_VS_MRL24_TREATED_MEF_DN
GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP, GSE19923_E2A_KO_VS_HEB_AND_E2A_KO_DP_THYMOCYTE_UP
GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_UP, GSE17721_LPS_VS_PAM3CSK4_2H_BMDC_UP
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
PID_HDAC_CLASSII_PATHWAY, PID_HDAC_CLASSII_PATHWAY
GSE360_CTRL_VS_L_MAJOR_DC_DN, GSE360_CTRL_VS_L_MAJOR_DC_DN
GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_UP, GSE25123_CTRL_VS_ROSIGLITAZONE_STIM_PPARG_KO_MACROPHAGE_UP
PID_HDAC_CLASSI_PATHWAY, PID_HDAC_CLASSI_PATHWAY
BROWNE_HCMV_INFECTION_12HR_UP, BROWNE_HCMV_INFECTION_12HR_UP
PRAMOONJAGO_SOX4_TARGETS_UP, PRAMOONJAGO_SOX4_TARGETS_UP
BURTON_ADIPOGENESIS_PEAK_AT_2HR, BURTON_ADIPOGENESIS_PEAK_AT_2HR
AMIT_EGF_RESPONSE_120_HELA, AMIT_EGF_RESPONSE_120_HELA
GSE22432_MULTIPOTENT_PROGENITOR_VS_PDC_UP, GSE22432_MULTIPOTENT_PROGENITOR_VS_PDC_UP
GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_DN, GSE1432_CTRL_VS_IFNG_1H_MICROGLIA_DN
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN
GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_0.5H_BMDC_DN
KEGG_P53_SIGNALING_PATHWAY, KEGG_P53_SIGNALING_PATHWAY
GO_NEGATIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT, GO_NEGATIVE_REGULATION_OF_NUCLEOCYTOPLASMIC_TRANSPORT
GO_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS, GO_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS
WANG_HCP_PROSTATE_CANCER, WANG_HCP_PROSTATE_CANCER
THEILGAARD_NEUTROPHIL_AT_SKIN_WOUND_UP, THEILGAARD_NEUTROPHIL_AT_SKIN_WOUND_UP
PID_HIV_NEF_PATHWAY, PID_HIV_NEF_PATHWAY
GO_PROTEIN_PHOSPHATASE_BINDING, GO_PROTEIN_PHOSPHATASE_BINDING
ELVIDGE_HIF1A_AND_HIF2A_TARGETS_DN, ELVIDGE_HIF1A_AND_HIF2A_TARGETS_DN
MODULE_237, MODULE_237
GO_NEGATIVE_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS, GO_NEGATIVE_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS
ZHANG_PROLIFERATING_VS_QUIESCENT, ZHANG_PROLIFERATING_VS_QUIESCENT
GO_REGULATION_OF_ATPASE_ACTIVITY, GO_REGULATION_OF_ATPASE_ACTIVITY
GO_SUMO_TRANSFERASE_ACTIVITY, GO_SUMO_TRANSFERASE_ACTIVITY
GSE22527_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_UP, GSE22527_ANTI_CD3_INVIVO_VS_UNTREATED_MOUSE_TREG_UP
BURTON_ADIPOGENESIS_1, BURTON_ADIPOGENESIS_1
BIOCARTA_DEATH_PATHWAY, BIOCARTA_DEATH_PATHWAY
GO_REGULATION_OF_MUSCLE_SYSTEM_PROCESS, GO_REGULATION_OF_MUSCLE_SYSTEM_PROCESS
GSE27786_LSK_VS_NEUTROPHIL_DN, GSE27786_LSK_VS_NEUTROPHIL_DN
GCM_MAPK10, GCM_MAPK10
GO_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL, GO_REGULATION_OF_MITOCHONDRIAL_MEMBRANE_POTENTIAL
GGCNRNWCTTYS_UNKNOWN, GGCNRNWCTTYS_UNKNOWN
GO_POSITIVE_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING, GO_POSITIVE_REGULATION_OF_NIK_NF_KAPPAB_SIGNALING
WANG_TARGETS_OF_MLL_CBP_FUSION_UP, WANG_TARGETS_OF_MLL_CBP_FUSION_UP
AMIT_EGF_RESPONSE_120_MCF10A, AMIT_EGF_RESPONSE_120_MCF10A
MODULE_521, MODULE_521
SMIRNOV_RESPONSE_TO_IR_2HR_DN, SMIRNOV_RESPONSE_TO_IR_2HR_DN
GO_STEROL_BIOSYNTHETIC_PROCESS, GO_STEROL_BIOSYNTHETIC_PROCESS
REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS, REACTOME_EXTRINSIC_PATHWAY_FOR_APOPTOSIS
GO_COPPER_ION_TRANSPORT, GO_COPPER_ION_TRANSPORT
KEGG_STEROID_BIOSYNTHESIS, KEGG_STEROID_BIOSYNTHESIS
GO_COPPER_ION_HOMEOSTASIS, GO_COPPER_ION_HOMEOSTASIS
GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS, GO_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_ENDOPLASMIC_RETICULUM_STRESS
REACTOME_CHOLESTEROL_BIOSYNTHESIS, REACTOME_CHOLESTEROL_BIOSYNTHESIS
BOGNI_TREATMENT_RELATED_MYELOID_LEUKEMIA_UP, BOGNI_TREATMENT_RELATED_MYELOID_LEUKEMIA_UP
GO_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH, GO_POSITIVE_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_CELL_DEATH
GO_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE, GO_STRUCTURAL_CONSTITUENT_OF_NUCLEAR_PORE
GO_LYMPHOCYTE_APOPTOTIC_PROCESS, GO_LYMPHOCYTE_APOPTOTIC_PROCESS
GO_NEGATIVE_REGULATION_OF_NF_KAPPAB_IMPORT_INTO_NUCLEUS, GO_NEGATIVE_REGULATION_OF_NF_KAPPAB_IMPORT_INTO_NUCLEUS
GO_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY, GO_NUCLEOTIDE_BINDING_DOMAIN_LEUCINE_RICH_REPEAT_CONTAINING_RECEPTOR_SIGNALING_PATHWAY
GO_SCHWANN_CELL_DIFFERENTIATION, GO_SCHWANN_CELL_DIFFERENTIATION
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS
GO_DEATH_RECEPTOR_ACTIVITY, GO_DEATH_RECEPTOR_ACTIVITY
GO_REGULATION_OF_NFAT_PROTEIN_IMPORT_INTO_NUCLEUS, GO_REGULATION_OF_NFAT_PROTEIN_IMPORT_INTO_NUCLEUS
GO_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA, GO_POSITIVE_REGULATION_OF_RELEASE_OF_CYTOCHROME_C_FROM_MITOCHONDRIA
YANG_BREAST_CANCER_ESR1_BULK_DN, YANG_BREAST_CANCER_ESR1_BULK_DN
GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_DN, GSE24972_MARGINAL_ZONE_BCELL_VS_FOLLICULAR_BCELL_DN
WEIGEL_OXIDATIVE_STRESS_BY_TBH_AND_H2O2, WEIGEL_OXIDATIVE_STRESS_BY_TBH_AND_H2O2
GO_RESPONSE_TO_OXYGEN_RADICAL, GO_RESPONSE_TO_OXYGEN_RADICAL
GO_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY, GO_REGULATION_OF_OXIDATIVE_STRESS_INDUCED_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
TCTGGAC_MIR198, TCTGGAC_MIR198
FARDIN_HYPOXIA_11, FARDIN_HYPOXIA_11
BIOCARTA_41BB_PATHWAY, BIOCARTA_41BB_PATHWAY
REACTOME_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY, REACTOME_POST_CHAPERONIN_TUBULIN_FOLDING_PATHWAY
GO_T_CELL_APOPTOTIC_PROCESS, GO_T_CELL_APOPTOTIC_PROCESS
KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_WS, KYNG_ENVIRONMENTAL_STRESS_RESPONSE_NOT_BY_UV_IN_WS
GO_CELLULAR_COPPER_ION_HOMEOSTASIS, GO_CELLULAR_COPPER_ION_HOMEOSTASIS
GO_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS, GO_NEGATIVE_REGULATION_OF_EXTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_VIA_DEATH_DOMAIN_RECEPTORS