

_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_6H_DN

GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN, GSE2770_TGFB_AND_IL4_ACT_VS_ACT_CD4_TCELL_6H_DN
REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE, REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE
GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_TGFB_AND_IL4_VS_IL12_TREATED_ACT_CD4_TCELL_48H_DN
GSE23695_CD57_POS_VS_NEG_NK_CELL_DN, GSE23695_CD57_POS_VS_NEG_NK_CELL_DN
HALLMARK_UV_RESPONSE_UP, HALLMARK_UV_RESPONSE_UP
GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_DN, GSE18791_CTRL_VS_NEWCASTLE_VIRUS_DC_12H_DN
GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP, GROSS_HYPOXIA_VIA_ELK3_AND_HIF1A_UP
GO_CELLULAR_RESPONSE_TO_UV, GO_CELLULAR_RESPONSE_TO_UV
MENSE_HYPOXIA_UP, MENSE_HYPOXIA_UP
GSE3982_CTRL_VS_LPS_48H_DC_DN, GSE3982_CTRL_VS_LPS_48H_DC_DN
ELVIDGE_HYPOXIA_BY_DMOG_UP, ELVIDGE_HYPOXIA_BY_DMOG_UP
GSE360_CTRL_VS_L_MAJOR_DC_DN, GSE360_CTRL_VS_L_MAJOR_DC_DN
GO_PROTEIN_PHOSPHATASE_BINDING, GO_PROTEIN_PHOSPHATASE_BINDING
GSE27786_LSK_VS_NEUTROPHIL_DN, GSE27786_LSK_VS_NEUTROPHIL_DN
WANG_HCP_PROSTATE_CANCER, WANG_HCP_PROSTATE_CANCER
GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN, GSE18791_UNSTIM_VS_NEWCATSLE_VIRUS_DC_1H_DN
FRASOR_RESPONSE_TO ESTRADIOL_DN, FRASOR_RESPONSE_TO ESTRADIOL_DN
AMIT_EGF_RESPONSE_120_HELA, AMIT_EGF_RESPONSE_120_HELA
GO_REGULATION_OF_MUSCLE_SYSTEM_PROCESS, GO_REGULATION_OF_MUSCLE_SYSTEM_PROCESS
GO_REGULATION_OF_ATPASE_ACTIVITY, GO_REGULATION_OF_ATPASE_ACTIVITY
AMIT_EGF_RESPONSE_120_MCF10A, AMIT_EGF_RESPONSE_120_MCF10A
GO_SCF_UBIQUITIN_LIGASE_COMPLEX, GO_SCF_UBIQUITIN_LIGASE_COMPLEX
ZHAN_MULTIPLE_MYELOMA_MS_DN, ZHAN_MULTIPLE_MYELOMA_MS_DN
GCM_MAPK10, GCM_MAPK10
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE
LABBE_WNT3A_TARGETS_DN, LABBE_WNT3A_TARGETS_DN
GO_MICROTUBULE_ANCHORING, GO_MICROTUBULE_ANCHORING
KEGG_PRION_DISEASES, KEGG_PRION_DISEASES
GO_POSITIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS, GO_POSITIVE_REGULATION_OF_RESPONSE_TO_OXIDATIVE_STRESS
TCTGGAC_MIR198, TCTGGAC_MIR198
SA_PROGRAMMED_CELL_DEATH, SA_PROGRAMMED_CELL_DEATH
GO_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS, GO_REGULATION_OF_TRANSCRIPTION_FACTOR_IMPORT_INTO_NUCLEUS
GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE_BY_P53_CLASS_MEDIATOR, GO_REGULATION_OF_INTRINSIC_APOPTOTIC_SIGNALING_PATHWAY_IN_RESPONSE_TO_DNA_DAMAGE_BY_P53_CLASS_MEDIATOR
KRIEG_HYPOXIA_VIA_KDM3A, KRIEG_HYPOXIA_VIA_KDM3A