

TANT_NOD2_TRANSDUCED_HEK293T_CELL_DN, GSE22611_UNSTIM_VS_2H_MDP_STIM_MUTANT_NOD2_TRANSDUCED_HEK293T_CELL_DN

GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_12H_VS_24H_F_TULARENSIS_LVS_NEUTROPHIL_UP
GSE14769_20MIN_VS_360MIN_LPS_BMDM_DN, GSE14769_20MIN_VS_360MIN_LPS_BMDM_DN
GSE40184_HEALTHY_VS_HCV_INFECTED_DONOR_PBMCDN, GSE40184_HEALTHY_VS_HCV_INFECTED_DONOR_PBMCDN
GSE4590_SMALL_VS_VPREB_POS_LARGE_PRE_BCELL_UP, GSE4590_SMALL_VS_VPREB_POS_LARGE_PRE_BCELL_UP
GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_4H_DN, GSE7831_UNSTIM_VS_INFLUENZA_STIM_PDC_4H_DN
GSE11924_TH1_VS_TH2_CD4_TCELL_UP, GSE11924_TH1_VS_TH2_CD4_TCELL_UP
GSE40068_CXCR5POS_BCL6POS_TFH_VS_CXCR5NEG_BCL6NEG_CD4_TCELL_DN, GSE40068_CXCR5POS_BCL6POS_TFH_VS_CXCR5NEG_BCL6NEG_CD4_TCELL_DN
TGAGATT_MIR216, TGAGATT_MIR216
ZHANG_TARGETS_OF_EWSR1_FLI1_FUSION, ZHANG_TARGETS_OF_EWSR1_FLI1_FUSION
GSE17721_CTRL_VS_LPS_8H_BMDC_DN, GSE17721_CTRL_VS_LPS_8H_BMDC_DN
KEGG_PHOSPHATIDYLOSITOL_SIGNALING_SYSTEM, KEGG_PHOSPHATIDYLOSITOL_SIGNALING_SYSTEM
GSE21379_TFH_VS_NON_TFH_CD4_TCELL_UP, GSE21379_TFH_VS_NON_TFH_CD4_TCELL_UP
CTACTGT_MIR199A, CTACTGT_MIR199A
GO_REGULATION_OF_MITOPHAGY, GO_REGULATION_OF_MITOPHAGY
GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY, GO_BETA_CATENIN_TCF_COMPLEX_ASSEMBLY
GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_IL12_CD8_TCELL_UP, GSE15930_NAIVE_VS_48H_IN_VITRO_STIM_IL12_CD8_TCELL_UP
MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_DN, MARIADASON_REGULATED_BY_HISTONE_ACETYLATION_DN
FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP, FONTAINE_FOLLICULAR_THYROID_ADENOMA_UP
GO_CARDIAC_CHAMBER_DEVELOPMENT, GO_CARDIAC_CHAMBER_DEVELOPMENT
GSE4984_UNTREATED_VS_LPS_TREATED_DC_UP, GSE4984_UNTREATED_VS_LPS_TREATED_DC_UP
GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_UP, GSE22589_HIV_VS_HIV_AND_SIV_INFECTED_DC_UP
GO_MRNA_CLEAVAGE_FACTOR_COMPLEX, GO_MRNA_CLEAVAGE_FACTOR_COMPLEX
MEIS1BHOXA9_01, MEIS1BHOXA9_01
GO_CARDIAC_CHAMBER_MORPHOGENESIS, GO_CARDIAC_CHAMBER_MORPHOGENESIS
GO_LIPID_PHOSPHORYLATION, GO_LIPID_PHOSPHORYLATION
EHLERS_ANEUPLOIDY_UP, EHLERS_ANEUPLOIDY_UP
GO_RNA_POLYADENYLATION, GO_RNA_POLYADENYLATION
GO_RESPONSE_TO_YEAST, GO_RESPONSE_TO_YEAST
GO_CARDIAC_VENTRICLE_DEVELOPMENT, GO_CARDIAC_VENTRICLE_DEVELOPMENT
BIOCARTA_P35ALZHEIMERS_PATHWAY, BIOCARTA_P35ALZHEIMERS_PATHWAY
KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION, KYNG_DNA_DAMAGE_BY_GAMMA_AND_UV_RADIATION
GO_REGULATION_OF_HORMONE_METABOLIC_PROCESS, GO_REGULATION_OF_HORMONE_METABOLIC_PROCESS
MORF_RUNX1, MORF_RUNX1
GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN, GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN
GO_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS, GO_ENDODEOXYRIBONUCLEASE_ACTIVITY_PRODUCING_5_PHOSPHOMONOESTERS
FIGUEROA_AML_METHYLATION_CLUSTER_7_DN, FIGUEROA_AML_METHYLATION_CLUSTER_7_DN
WATTEL_AUTONOMOUS_THYROID_ADENOMA_UP, WATTEL_AUTONOMOUS_THYROID_ADENOMA_UP
GO_CELL_GROWTH, GO_CELL_GROWTH
GO_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR_RESPONSE_TO_GLUCOSE_STIMULUS, GO_REGULATION_OF_INSULIN_SECRETION_INVOLVED_IN_CELLULAR
GO_MATING, GO_MATING
ZHAN_MULTIPLE_MYELOMA_MF_DN, ZHAN_MULTIPLE_MYELOMA_MF_DN
GO_CARDIAC_VENTRICLE_MORPHOGENESIS, GO_CARDIAC_VENTRICLE_MORPHOGENESIS
GO_EMBRYONIC_HEMOPOIESIS, GO_EMBRYONIC_HEMOPOIESIS
LANDIS_ERBB2_BREAST_TUMORS_65_DN, LANDIS_ERBB2_BREAST_TUMORS_65_DN
REACTOME_CD28_DEPENDENT_VAV1_PATHWAY, REACTOME_CD28_DEPENDENT_VAV1_PATHWAY
GO TRABECULA_MORPHOGENESIS, GO TRABECULA_MORPHOGENESIS