

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
GSE32901_NAIVE_VS_TH17_ENRICHED_CD4_TCELL_DN, GSE32901_NAIVE_VS_TH17_ENRICHED_CD4_TCELL_DN
GSE22935_WT_VS_MYPD88_KO_MACROPHAGE_24H_MBOVIS_BCG_STIM_DN, GSE22935_WT_VS_MYPD88_KO_MACROPHAGE_24H_MBOVIS_BCG_STIM_DN
GSE14769_20MIN_VS_360MIN_LPS_BMDM_DN, GSE14769_20MIN_VS_360MIN_LPS_BMDM_DN
GSE17721_0.5H_VS_12H_LPS_BMDC_DN, GSE17721_0.5H_VS_12H_LPS_BMDC_DN
GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_DN, GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_DN
GSE40068_CXCR5POS_BCL6POS_TFH_VS_CXCR5NEG_BCL6NEG_CD4_TCELL_DN, GSE40068_CXCR5POS_BCL6POS_TFH_VS_CXCR5NEG_BCL6NEG_CD4_TCELL_DN
GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN, GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN
GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17_POLARIZING_CONDITIONS_DN, GSE27241_CTRL_VS_DIGOXIN_TREATED_RORGT_KO_CD4_TCELL_IN_TH17
GO_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS, GO_PHOSPHATIDYLINOSITOL_METABOLIC_PROCESS
GO_INOSITOL_PHOSPHATE_METABOLIC_PROCESS, GO_INOSITOL_PHOSPHATE_METABOLIC_PROCESS
COMP1_01, COMP1_01
SASSON_RESPONSE_TO_FORSKOLIN_DN, SASSON_RESPONSE_TO_FORSKOLIN_DN
HOLLEMAN_VINCRISTINE_RESISTANCE_ALL_UP, HOLLEMAN_VINCRISTINE_RESISTANCE_ALL_UP
GO_NEGATIVE_REGULATION_OF_TOR_SIGNALING, GO_NEGATIVE_REGULATION_OF_TOR_SIGNALING
GO_HSP90_PROTEIN_BINDING, GO_HSP90_PROTEIN_BINDING
BIOCARTA_AKAP13_PATHWAY, BIOCARTA_AKAP13_PATHWAY
GSE3982_MAC_VS_NKCELL_DN, GSE3982_MAC_VS_NKCELL_DN
GO_LIPID_PHOSPHORYLATION, GO_LIPID_PHOSPHORYLATION
GO_CARDIAC_VENTRICLE_DEVELOPMENT, GO_CARDIAC_VENTRICLE_DEVELOPMENT
GO_LYSOPHOSPHOLIPID_ACYLTRANSFERASE_ACTIVITY, GO_LYSOPHOSPHOLIPID_ACYLTRANSFERASE_ACTIVITY
REACTOME_CD28_DEPENDENT_VAV1_PATHWAY, REACTOME_CD28_DEPENDENT_VAV1_PATHWAY
SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_UP, SCHAEFFER_PROSTATE_DEVELOPMENT_AND_CANCER_BOX1_UP
YTAATTAA_LHX3_01, YTAATTAA_LHX3_01
GO_REGULATION_OF_RECEPTOR_ACTIVITY, GO_REGULATION_OF_RECEPTOR_ACTIVITY
chr4q26, chr4q26
KUNINGER_IGF1_VS_PDGFB_TARGETS_DN, KUNINGER_IGF1_VS_PDGFB_TARGETS_DN
GO_RESPONSE_TO_YEAST, GO_RESPONSE_TO_YEAST
GO_ESTABLISHMENT_OF_CELL_POLARITY, GO_ESTABLISHMENT_OF_CELL_POLARITY
GO_REGULATION_OF_HORMONE_METABOLIC_PROCESS, GO_REGULATION_OF_HORMONE_METABOLIC_PROCESS
FONTAINE_PAPILLARY_THYROID_CARCINOMA_DN, FONTAINE_PAPILLARY_THYROID_CARCINOMA_DN
GO_CELLULAR_RESPONSE_TO_ELECTRICAL_STIMULUS, GO_CELLULAR_RESPONSE_TO_ELECTRICAL_STIMULUS
KEGG_CIRCADIAN_RHYTHM_MAMMAL, KEGG_CIRCADIAN_RHYTHM_MAMMAL