

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
- GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_DN, GSE26928_NAIVE_VS_CXCR5_POS_CD4_TCELL_DN
- GSE360_CTRL_VS_L_MAJOR_DC_UP, GSE360_CTRL_VS_L_MAJOR_DC_UP
- GSE3039_CD4_TCELL_VS_B1_BCELL_UP, GSE3039_CD4_TCELL_VS_B1_BCELL_UP
- GOBP_GOLGI_ORGANIZATION, GOBP_GOLGI_ORGANIZATION
- GSE40184_HEALTHY_VS_HCV_INFECTED_DONOR_PBMCDN, GSE40184_HEALTHY_VS_HCV_INFECTED_DONOR_PBMCDN
- GSE29949_CD8_NEG_DC_SPLEEN_VS_DC_BRAIN_DN, GSE29949_CD8_NEG_DC_SPLEEN_VS_DC_BRAIN_DN
- GSE22935_WT_VS_MYPD88_KO_MACROPHAGE_12H_MBOVIS_BCG_STIM_UP, GSE22935_WT_VS_MYPD88_KO_MACROPHAGE_12H_MBOVIS_BCG_STIM_UP
- GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_MOUSE_LUNG_DC_UP, GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_MOUSE_LUNG_DC_UP
- GOCC_HISTONE_METHYLTRANSFERASE_COMPLEX, GOCC_HISTONE_METHYLTRANSFERASE_COMPLEX
- GSE22589_HEALTHY_VS_SIV_INFECTED_DC_DN, GSE22589_HEALTHY_VS_SIV_INFECTED_DC_DN
- KMCATNNWGGA_UNKNOWN, KMCATNNWGGA_UNKNOWN
- PCGF6_TARGET_GENES, PCGF6_TARGET_GENES
- GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN, GSE2585_THYMIC_MACROPHAGE_VS_MTEC_DN
- MIR6775_3P, MIR6775_3P
- MIR4288, MIR4288
- GSE18281_CORTEX_VS_MEDULLA_THYMUS_UP, GSE18281_CORTEX_VS_MEDULLA_THYMUS_UP
- MIR3173_5P, MIR3173_5P
- MIR6799_3P, MIR6799_3P
- HP_PULMONARY_ARTERY_STENOSIS, HP_PULMONARY_ARTERY_STENOSIS
- MIR7158_3P, MIR7158_3P
- PRDM12_TARGET_GENES, PRDM12_TARGET_GENES
- GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_CHEMOTAXIS, GOBP_POSITIVE_REGULATION_OF_GRANULOCYTE_CHEMOTAXIS
- HP_PATELLAR_DISLOCATION, HP_PATELLAR_DISLOCATION
- HP_CHILDHOOD_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT, HP_CHILDHOOD_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT
- MIR3157_3P, MIR3157_3P
- EHLERS_ANEUPLOIDY_UP, EHLERS_ANEUPLOIDY_UP
- GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS, GOBP_PHOSPHATIDIC_ACID_METABOLIC_PROCESS
- MIR151A_5P_MIR151B, MIR151A_5P_MIR151B
- GCGCCTT_MIR525_MIR524, GCGCCTT_MIR525_MIR524
- KEGG_CIRCADIAN_RHYTHM_MAMMAL, KEGG_CIRCADIAN_RHYTHM_MAMMAL
- HP_CHYLOTHORAX, HP_CHYLOTHORAX
- HARALAMBIEVA_PBMCFUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID_GENE_MODELS_PRED_PEAK_B_CELL_ELISPOT_RESP_NEGATIVE, HARALAMBIEVA_PBMCFUARIX_AGE_50_74YO_CORR_WITH_28D_MEM_B_CELL_RESPONSE_AT_28DY_LATE_GENE_EXPR_INDIVID_GENE_MODELS_PRED
- TAKAYAMA_BOUND_BY_AR, TAKAYAMA_BOUND_BY_AR
- GOBP_PRIMARY_ALCOHOL_BIOSYNTHETIC_PROCESS, GOBP_PRIMARY_ALCOHOL_BIOSYNTHETIC_PROCESS