

F_TULARENSIS_LVS_NEUTROPHIL_UP, GSE37416_0H_VS_3H_F_TULARENSIS_LVS_NEUTROPHIL_UP

GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_NONATOPIC_PATIENT_DN, GSE14908_RESTING_VS_HDM_STIM_CD4_TCELL_NONATOPIC_PATIENT_DN
GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_TGFB_AND_IL4_VS_TGFB_AND_IL12_TREATED_ACT_CD4_TCELL_48H_DN
GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP, GSE5503_MLN_DC_VS_SPLEEN_DC_ACTIVATED_ALLOGENIC_TCELL_UP
GSE17721_PAM3CSK4_VS_GADIQUIMOD_6H_BMDC_DN, GSE17721_PAM3CSK4_VS_GADIQUIMOD_6H_BMDC_DN
REACTOME_RAB_REGULATION_OF_TRAFFICKING, REACTOME_RAB_REGULATION_OF_TRAFFICKING
GOCC_SPECIFIC_GRANULE_MEMBRANE, GOCC_SPECIFIC_GRANULE_MEMBRANE
GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELL_UP, GSE3039_CD4_TCELL_VS_ALPHABETA_CD8_TCELL_UP
GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP, GSE29164_DAY3_VS_DAY7_CD8_TCELL_TREATED_MELANOMA_UP
GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_UP, GSE4984_UNTREATED_VS_VEHICLE_CTRL_TREATED_DC_UP
GSE17721_CTRL_VS_CPG_24H_BMDC_UP, GSE17721_CTRL_VS_CPG_24H_BMDC_UP
GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP, GSE34156_TLR1_TLR2_LIGAND_VS_NOD2_AND_TLR1_TLR2_LIGAND_6H_TREATED_MONOCYTE_UP
BIOCARTA_HIVNEF_PATHWAY, BIOCARTA_HIVNEF_PATHWAY
CHANDRAN_METASTASIS_TOP50_DN, CHANDRAN_METASTASIS_TOP50_DN
GSE35543_IN_VIVO_NTREG_VS_IN_VITRO_ITREG_DN, GSE35543_IN_VIVO_NTREG_VS_IN_VITRO_ITREG_DN
GOBP_NEGATIVE_REGULATION_OF_ORGANELLE_ASSEMBLY, GOBP_NEGATIVE_REGULATION_OF_ORGANELLE_ASSEMBLY
REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS, REACTOME_GLYCEROPHOSPHOLIPID_BIOSYNTHESIS
MIR10397_5P, MIR10397_5P
MIR5581_3P, MIR5581_3P
GSE32901_NAIVE_VS_TH1_CD4_TCELL_UP, GSE32901_NAIVE_VS_TH1_CD4_TCELL_UP
MIR3152_5P, MIR3152_5P
MODULE_486, MODULE_486
GOCC_ORGANELLE_MEMBRANE_CONTACT_SITE, GOCC_ORGANELLE_MEMBRANE_CONTACT_SITE
MIR504_5P, MIR504_5P
MIR4725_5P, MIR4725_5P
GOBP_RESPONSE_TO_ANTIBIOTIC, GOBP_RESPONSE_TO_ANTIBIOTIC
MIR4329, MIR4329
HP_CHRONIC_LEUKEMIA, HP_CHRONIC_LEUKEMIA
GOCC_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK, GOCC_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK
GOBP_INTERMEMBRANE_LIPID_TRANSFER, GOBP_INTERMEMBRANE_LIPID_TRANSFER
GOMF_LIPID_TRANSFER_ACTIVITY, GOMF_LIPID_TRANSFER_ACTIVITY
MIR6751_5P, MIR6751_5P
MIR12121, MIR12121
GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS, GOBP_POSITIVE_REGULATION_OF_DENDRITIC_SPINE_MORPHOGENESIS
GOMF_METALLOEXOPEPTIDASE_ACTIVITY, GOMF_METALLOEXOPEPTIDASE_ACTIVITY
GOBP_PHOSPHATIDYLSERINE_ACYL_CHAIN_REMODELING, GOBP_PHOSPHATIDYLSERINE_ACYL_CHAIN_REMODELING
REACTOME_ACYL_CHAIN_REMODELLING_OF_PS, REACTOME_ACYL_CHAIN_REMODELLING_OF_PS
MIR625_3P, MIR625_3P
EVI1_03, EVI1_03
MIR323A_5P, MIR323A_5P
GOBP_REGULATION_OF_DNA_CATABOLIC_PROCESS, GOBP_REGULATION_OF_DNA_CATABOLIC_PROCESS
REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOINOSITIDE_3_KINASE_PI3K, REACTOME_ERYTHROPOIETIN_ACTIVATES_PHOSPHOINOSITIDE_3_KINASE_PI3K
GOBP_POSITIVE_REGULATION_OF_CHROMATIN_BINDING, GOBP_POSITIVE_REGULATION_OF_CHROMATIN_BINDING
MIR503_3P, MIR503_3P
GOCC_SMN_COMPLEX, GOCC_SMN_COMPLEX
HP_ABNORMALITY_OF_THE_DISTAL_PHALANGES_OF_THE_TOES, HP_ABNORMALITY_OF_THE_DISTAL_PHALANGES_OF_THE_TOES
NR5A1_TARGET_GENES, NR5A1_TARGET_GENES