

GSE11864_CS_F1_IFNG_PAM3CYS_IN_MAC_UP, GSE11864_CS_F1_IFNG_PAM3CYS_IN_MAC_UP

GSE2706_UNSTIM_VS_8H_LPS_DC_UP, GSE2706_UNSTIM_VS_8H_LPS_DC_UP
GSE11864_UNTREATED_VS_CS_F1_IFNG_PAM3CYS_IN_MAC_UP, GSE11864_UNTREATED_VS_CS_F1_IFNG_PAM3CYS_IN_MAC_UP
GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN, GSE2770_TGFB_AND_IL4_VS_IL4_TREATED_ACT_CD4_TCELL_48H_DN
GSE11864_CS_F1_VS_CS_F1_IFNG_PAM3CYS_IN_MAC_UP, GSE11864_CS_F1_VS_CS_F1_IFNG_PAM3CYS_IN_MAC_UP
GSE26495_NAIVE_VS_PD1LOW_CD8_TCELL_DN, GSE26495_NAIVE_VS_PD1LOW_CD8_TCELL_DN
GSE11057_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN, GSE11057_NAIVE_VS_EFF_MEMORY_CD4_TCELL_DN
GSE42724_NAIVE_BCELL_VS_PLASMABLAST_DN, GSE42724_NAIVE_BCELL_VS_PLASMABLAST_DN
GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_DN, GSE22611_NOD2_VS_CTRL_TRANSDUCED_HEK293T_CELL_DN
GSE2706_2H_VS_8H_LPS_STIM_DC_UP, GSE2706_2H_VS_8H_LPS_STIM_DC_UP
GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN, GSE15324_ELF4_KO_VS_WT_ACTIVATED_CD8_TCELL_DN
GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN, GSE1460_CD4_THYMOCYTE_VS_NAIVE_CD4_TCELL_CORD_BLOOD_DN
THAKAR_PBMK_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_28DY_DN, THAKAR_PBMK_INACTIVATED_INFLUENZA_AGE_21_30YO_NONRESPONDER_28DY_DN
MODULE_430, MODULE_430
GOBP_NEGATIVE_REGULATION_OF_MAPK_CASCADE, GOBP_NEGATIVE_REGULATION_OF_MAPK_CASCADE
GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION, GOBP_REGULATION_OF_RHO_PROTEIN_SIGNAL_TRANSDUCTION
HP_IMPAIRED_VIBRATION_SENSATION_IN_THE_LOWER_LIMBS, HP_IMPAIRED_VIBRATION_SENSATION_IN_THE_LOWER_LIMBS
ZNF510_TARGET_GENES, ZNF510_TARGET_GENES
GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN, GSE37301_LYMPHOID_PRIMED_MPP_VS_PRO_BCELL_DN
HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_1DY_DN, HOEK_NEUTROPHIL_2011_2012_TIV_ADULT_1DY_DN
GOBP_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY, GOBP_REGULATION_OF_G_PROTEIN_COUPLED_RECEPTOR_SIGNALING_PATHWAY
MIR1298_3P, MIR1298_3P
GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING, GOBP_POSITIVE_REGULATION_OF_PROTEIN_BINDING
HP_ABNORMALITY_OF_THE_TALUS, HP_ABNORMALITY_OF_THE_TALUS
MIR6857_3P, MIR6857_3P
ZNF433_TARGET_GENES, ZNF433_TARGET_GENES
GOMF_PHOSPHOLIPASE_ACTIVITY, GOMF_PHOSPHOLIPASE_ACTIVITY
MIR4703_3P, MIR4703_3P
MIR599, MIR599
GOBP_ACTIN_CYTOSKELETON_REORGANIZATION, GOBP_ACTIN_CYTOSKELETON_REORGANIZATION
MIR6089, MIR6089
GOCC_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK, GOCC_ENDOPLASMIC_RETICULUM_TUBULAR_NETWORK
MODULE_247, MODULE_247
FOXD3_TARGET_GENES, FOXD3_TARGET_GENES
WP_EICOSANOID_METABOLISM_VIA_LIPO_OXYGENASES_LOX, WP_EICOSANOID_METABOLISM_VIA_LIPO_OXYGENASES_LOX
MIR3681_5P, MIR3681_5P
GOCC_VOLTAGE_GATED_CALCIUM_CHANNEL_COMPLEX, GOCC_VOLTAGE_GATED_CALCIUM_CHANNEL_COMPLEX
ZNF555_TARGET_GENES, ZNF555_TARGET_GENES
GOCC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_COMPLEX, GOCC_GUANYL_NUCLEOTIDE_EXCHANGE_FACTOR_COMPLEX
REACTOME_MET_ACTIVATES_RAP1_AND_RAC1, REACTOME_MET_ACTIVATES_RAP1_AND_RAC1
TAVAZOIE_METASTASIS, TAVAZOIE_METASTASIS
MIR6849_5P, MIR6849_5P
RUAN_RESPONSE_TO_TROGLITAZONE_DN, RUAN_RESPONSE_TO_TROGLITAZONE_DN
GOBP_CELLULAR_MODIFIED_AMINO_ACID_CATABOLIC_PROCESS, GOBP_CELLULAR_MODIFIED_AMINO_ACID_CATABOLIC_PROCESS
GOBP_PRESYNAPTIC_ACTIVE_ZONE_ORGANIZATION, GOBP_PRESYNAPTIC_ACTIVE_ZONE_ORGANIZATION
GSE13173_UNTREATED_VS_IL12_TREATED_ACT_CD8_TCELL_UP, GSE13173_UNTREATED_VS_IL12_TREATED_ACT_CD8_TCELL_UP
LEE_LIVER_CANCER_MYC_DN, LEE_LIVER_CANCER_MYC_DN
GOBP_ZYMOGEN_INHIBITION, GOBP_ZYMOGEN_INHIBITION
GOMF_CHLORIDE_CHANNEL_REGULATOR_ACTIVITY, GOMF_CHLORIDE_CHANNEL_REGULATOR_ACTIVITY
HP_ABNORMALITY_OF_LENS_SHAPE, HP_ABNORMALITY_OF_LENS_SHAPE
chr3q27, chr3q27
HP_ADULT_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT, HP_ADULT_ONSET_SENSORINEURAL_HEARING_IMPAIRMENT