

B_MALAYI_HIGH_DOSE_DC_DN, GSE360_L_DONOVANI_VS_B_MALAYI_HIGH_DOSE_DC_DN

GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP, GSE360_CTRL_VS_M_TUBERCULOSIS_DC_UP
GSE28726_NAIVE_VS_ACTIVATED_CD4_TCELL_UP, GSE28726_NAIVE_VS_ACTIVATED_CD4_TCELL_UP
GSE360_CTRL_VS_M_TUBERCULOSIS_MAC_UP, GSE360_CTRL_VS_M_TUBERCULOSIS_MAC_UP
GSE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_MAC_DN, GSE360_T_GONDII_VS_B_MALAYI_LOW_DOSE_MAC_DN
GSE19401_PAM2CSK4_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM_FOLLICULAR_DC_DN, GSE19401_PAM2CSK4_VS_RETINOIC_ACID_AND_PAM2CSK4_STIM
GSE360_L_MAJOR_VS_B_MALAYI_HIGH_DOSE_DC_DN, GSE360_L_MAJOR_VS_B_MALAYI_HIGH_DOSE_DC_DN
GSE360_CTRL_VS_L_MAJOR_DC_UP, GSE360_CTRL_VS_L_MAJOR_DC_UP
GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_UP, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_MAC_UP
GSE360_DC_VS_MAC_DN, GSE360_DC_VS_MAC_DN
WAMUNYOKOLI_OVARIAN_CANCER_LMP_DN, WAMUNYOKOLI_OVARIAN_CANCER_LMP_DN
GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_MAC_DN, GSE360_L_DONOVANI_VS_B_MALAYI_LOW_DOSE_MAC_DN
GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN, GSE21927_SPLEEN_MONOCYTE_VS_GMCSF_GCSF_BONE_MARROW_DN
GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_MAC_DN, GSE360_LOW_DOSE_B_MALAYI_VS_M_TUBERCULOSIS_MAC_DN
MIR1205, MIR1205
GSE24574_BCL6_HIGH_TFH_VS_NAIVE_CD4_TCELL_UP, GSE24574_BCL6_HIGH_TFH_VS_NAIVE_CD4_TCELL_UP
GSE28726_NAIVE_VS_ACTIVATED_VA24NEG_NKTCELL_DN, GSE28726_NAIVE_VS_ACTIVATED_VA24NEG_NKTCELL_DN
GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_DN, GSE360_CTRL_VS_B_MALAYI_HIGH_DOSE_DC_DN
HP_DEVIATION_OF_TOES, HP_DEVIATION_OF_TOES
HP_BLEPHAROPHIMOSIS, HP_BLEPHAROPHIMOSIS
HP_APLASIA_HYPOPLASIA_INVOLVING_BONES_OF_THE_THORAX, HP_APLASIA_HYPOPLASIA_INVOLVING_BONES_OF_THE_THORAX
GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_DC_DN, GSE360_CTRL_VS_B_MALAYI_LOW_DOSE_DC_DN
PTEN_DN.V2_DN, PTEN_DN.V2_DN
KEGG_SPHINGOLIPID_METABOLISM, KEGG_SPHINGOLIPID_METABOLISM
GSE2706_R848_VS_LPS_8H_STIM_DC_UP, GSE2706_R848_VS_LPS_8H_STIM_DC_UP
GSE4142_NAIVE_VS_MEMORY_BCELL_DN, GSE4142_NAIVE_VS_MEMORY_BCELL_DN
MIR6760_5P, MIR6760_5P
GOCC_CIS_GOLGI_NETWORK, GOCC_CIS_GOLGI_NETWORK
REACTOME_RND1_GTPASE_CYCLE, REACTOME_RND1_GTPASE_CYCLE
HP_ABNORMAL_CORONARY_ARTERY_MORPHOLOGY, HP_ABNORMAL_CORONARY_ARTERY_MORPHOLOGY
HP_BROAD_PHALANX_OF_THE_TOES, HP_BROAD_PHALANX_OF_THE_TOES
RUNNE_GENDER_EFFECT_UP, RUNNE_GENDER_EFFECT_UP
WP_FARNESOID_X_RECEPTOR_PATHWAY, WP_FARNESOID_X_RECEPTOR_PATHWAY
GOCC_INTEGRAL_COMPONENT_OF_PRESYNAPTIC_ACTIVE_ZONE_MEMBRANE, GOCC_INTEGRAL_COMPONENT_OF_PRESYNAPTIC_ACTIVE_ZONE_MEM
MIR10396A_5P, MIR10396A_5P
MIR744_5P, MIR744_5P
GOCC_INTRINSIC_COMPONENT_OF_PRESYNAPTIC_ACTIVE_ZONE_MEMBRANE, GOCC_INTRINSIC_COMPONENT_OF_PRESYNAPTIC_ACTIVE_ZONE_MEM
REACTOME_TRAF6_MEDIATED_INDUCATION_OF_TAK1_COMPLEX_WITHIN_TLR4_COMPLEX, REACTOME_TRAF6_MEDIATED_INDUCATION_OF_TAK1_COMP
HP_EEG_WITH_CENTROTEMPORAL_FOCAL_SPIKE_WAVES, HP_EEG_WITH_CENTROTEMPORAL_FOCAL_SPIKE_WAVES