

1 VS 120MIN_LPS_BMDM_UP, GSE14769_UNSTIM_VS_120MIN_LPS_BMDM_UP

GSE26343_WT_VS_NFAT5_KO_MACROPHAGE_UP, GSE26343_WT_VS_NFAT5_KO_MACROPHAGE_UP
GSE40273_EOS_KO_VS_WT_TREG_DN, GSE40273_EOS_KO_VS_WT_TREG_DN
GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SPLEEN_6H_UP, GSE7768_OVA_ALONE_VS_OVA_WITH_LPS_IMMUNIZED_MOUSE_WHOLE_SP
GSE26343_WT_VS_NFAT5_KO_MACROPHAGE_LPS_STIM_UP, GSE26343_WT_VS_NFAT5_KO_MACROPHAGE_LPS_STIM_UP
GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_UP, GSE411_UNSTIM_VS_400MIN_IL6_STIM_MACROPHAGE_UP
GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_WITH_IFNA_STIM_90MIN_UP, GSE40666_NAIVE_VS_EFFECTOR_CD8_TCELL_WITH_IFNA_STIM_90MIN_UP
GSE4142_GC_BCELL_VS_MEMORY_BCELL_UP, GSE4142_GC_BCELL_VS_MEMORY_BCELL_UP
REACTOME_PI_METABOLISM, REACTOME_PI_METABOLISM
SIPA1_TARGET_GENES, SIPA1_TARGET_GENES
GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_UP, GSE36078_UNTREATED_VS_AD5_T425A_HEXON_INF_IL1R_KO_MOUSE_LUNG_DC_U
GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4HIGH_SORTED_BCELL_UP, GSE46606_UNSTIM_VS_CD40L_IL2_IL5_1DAY_STIMULATED_IRF4HIGH_SORTED
GSE32901_TH17_EMRICHD_VS_TH17_NEG_CD4_TCELL_UP, GSE32901_TH17_EMRICHD_VS_TH17_NEG_CD4_TCELL_UP
GOBP_REGULATION_OF_TORC1_SIGNALING, GOBP_REGULATION_OF_TORC1_SIGNALING
GOBP_PYRIMIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS, GOBP_PYRIMIDINE_CONTAINING_COMPOUND_CATABOLIC_PROCESS
GOBP_REGULATION_OF_EMBRYONIC_DEVELOPMENT, GOBP_REGULATION_OF_EMBRYONIC_DEVELOPMENT
GOCC_PALMITOYLTRANSFERASE_COMPLEX, GOCC_PALMITOYLTRANSFERASE_COMPLEX
GOCC_ENDOPLASMIC_RETICULUM_PALMITOYLTRANSFERASE_COMPLEX, GOCC_ENDOPLASMIC_RETICULUM_PALMITOYLTRANSFERASE_COMPLEX
GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION, GOBP_BASE_EXCISION_REPAIR_AP_SITE_FORMATION
REACTOME_PEXOPHAGY, REACTOME_PEXOPHAGY
GOBP_POSITIVE_REGULATION_OF_TORC1_SIGNALING, GOBP_POSITIVE_REGULATION_OF_TORC1_SIGNALING
GOBP_RESPONSE_TO_ACIDIC_PH, GOBP_RESPONSE_TO_ACIDIC_PH
MIR604, MIR604
GOBP_REGULATION_OF_FIBRINOLYSIS, GOBP_REGULATION_OF_FIBRINOLYSIS
HP_BLEPHAROCHALASIS, HP_BLEPHAROCHALASIS