

L10_STIM_IL10_KO_MACROPHAGE_10MIN_DN, GSE9509_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_10MIN_DN

GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN, GSE46606_IRF4MID_VS_WT_CD40L_IL2_IL5_DAY3_STIMULATED_BCELL_DN
GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_DN, GSE10273_HIGH_VS_LOW_IL7_TREATED_IRF4_8_NULL_PRE_BCELL_DN
GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_UP, GSE9650_EFFECTOR_VS_EXHAUSTED_CD8_TCELL_UP
GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP, GSE5463_CTRL_VS_DEXAMETHASONE_TREATED_THYMOCYTE_UP
GSE12198_CTRL_VS_LOW_IL2_STIM_NK_CELL_UP, GSE12198_CTRL_VS_LOW_IL2_STIM_NK_CELL_UP
GSE35685_CD34POS_CD10NEG_CD62LPOS_VS_CD34POS_CD10POS_BONE_MARROW_UP, GSE35685_CD34POS_CD10NEG_CD62LPOS_VS_CD34POS_CD10POS_BONE_MARROW_UP
GSE41867_DAY6_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN, GSE41867_DAY6_VS_DAY15_LCMV_ARMSTRONG_EFFECTOR_CD8_TCELL_DN
GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP, GSE44649_NAIVE_VS_ACTIVATED_CD8_TCELL_UP
GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_UP, GSE17721_CTRL_VS_GARDIQUIMOD_6H_BMDC_UP
GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_4H_ACT_CD4_TCELL_UP, GSE17974_IL4_AND_ANTI_IL12_VS_UNTREATED_4H_ACT_CD4_TCELL_UP
GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_GATA1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE37301_MULTIPOTENT_PROGENITOR_VS_COMMON_LYMPHOID_PROGENITOR_UP, GSE37301_MULTIPOTENT_PROGENITOR_VS_COMMON_LYMPHOID_PROGENITOR_UP
GSE17721_CTRL_VS_POLYIC_0.5H_BMDC_UP, GSE17721_CTRL_VS_POLYIC_0.5H_BMDC_UP
GO_LOCALIZATION_WITHIN_MEMBRANE, GO_LOCALIZATION_WITHIN_MEMBRANE
GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP, GSE33425_CD161_HIGH_VS_NEG_CD8_TCELL_UP
SETLUR_PROSTATE_CANCER_TMPRS2_ERG_FUSION_UP, SETLUR_PROSTATE_CANCER_TMPRS2_ERG_FUSION_UP
GSE28737_WT_VS_BCL6_HET_FOLLICULAR_BCELL_UP, GSE28737_WT_VS_BCL6_HET_FOLLICULAR_BCELL_UP
MODULE_430, MODULE_430
CHIARETTI_ACUTE_LYMPHOBLASTIC_LEUKEMIA_ZAP70, CHIARETTI_ACUTE_LYMPHOBLASTIC_LEUKEMIA_ZAP70
GO_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA, GO_RESPONSE_TO_TRANSFORMING_GROWTH_FACTOR_BETA
GO_RECEPTOR_CLUSTERING, GO_RECEPTOR_CLUSTERING
BIOCARTA_D4GDI_PATHWAY, BIOCARTA_D4GDI_PATHWAY
XU_RESPONSE_TO_TRETINOIN_AND_NSC682994_UP, XU_RESPONSE_TO_TRETINOIN_AND_NSC682994_UP
OXFORD_RALA_OR_RALB_TARGETS_DN, OXFORD_RALA_OR_RALB_TARGETS_DN
GO_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_BIOSYNTHETIC_PROCESS, GO_POSITIVE_REGULATION_OF_INTERFERON_GAMMA_BIOSYNTHETIC_PROCESS
GO_REGULATION_OF_MUSCLE_ADAPTATION, GO_REGULATION_OF_MUSCLE_ADAPTATION
GO_POSTSYNAPTIC_MEMBRANE_ORGANIZATION, GO_POSTSYNAPTIC_MEMBRANE_ORGANIZATION
REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION, REACTOME_TRAF6_MEDIATED_IRF7_ACTIVATION
REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY, REACTOME_TRAF3_DEPENDENT_IRF_ACTIVATION_PATHWAY
GO_REGULATION_OF_INTERFERON_GAMMA_BIOSYNTHETIC_PROCESS, GO_REGULATION_OF_INTERFERON_GAMMA_BIOSYNTHETIC_PROCESS
GO_AMINO_SUGAR_CATABOLIC_PROCESS, GO_AMINO_SUGAR_CATABOLIC_PROCESS
GO_RESPONSE_TO_EXOGENOUS_DSRNA, GO_RESPONSE_TO_EXOGENOUS_DSRNA
chr6q25, chr6q25