

_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_IRF4_VS_FOXP3_AND_IRF4_TRANSDUCED_ACTIVATED_CD4_TCELL_UP

GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_DN, GSE9650_EXHAUSTED_VS_MEMORY_CD8_TCELL_DN
GSE40274_FOXP3_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN, GSE40274_FOXP3_VS_FOXP3_AND_SATB1_TRANSDUCED_ACTIVATED_CD4_TCELL_DN
GSE45365_WT_VS_IFNAR_KO_CD11B_DC_UP, GSE45365_WT_VS_IFNAR_KO_CD11B_DC_UP
GSE16522_MEMORY_VS_NAIVE_CD8_TCELL_UP, GSE16522_MEMORY_VS_NAIVE_CD8_TCELL_UP
GSE17721_POLYIC_VS_GARDIQUIMOD_8H_BMDC_UP, GSE17721_POLYIC_VS_GARDIQUIMOD_8H_BMDC_UP
GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_DN, GSE1460_INTRATHYMIC_T_PROGENITOR_VS_DP_THYMOCYTE_DN
GSE17721_PAM3CSK4_VS_CPG_4H_BMDC_DN, GSE17721_PAM3CSK4_VS_CPG_4H_BMDC_DN
GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_NFATC1_KO_16H_UP, GSE21063_CTRL_VS_ANTI_IGM_STIM_BCELL_NFATC1_KO_16H_UP
GO_LAMELLIPODIUM, GO_LAMELLIPODIUM
GSE15930_NAIVE_VS_24H_IN_VITRO_STIM_CD8_TCELL_UP, GSE15930_NAIVE_VS_24H_IN_VITRO_STIM_CD8_TCELL_UP
BOYALT_LIVER_CANCER_SUBCLASS_G1_DN, BOYALT_LIVER_CANCER_SUBCLASS_G1_DN
GO_POSITIVE_REGULATION_OF_CYTOSKELETON_ORGANIZATION, GO_POSITIVE_REGULATION_OF_CYTOSKELETON_ORGANIZATION
KEGG_ADHERENS_JUNCTION, KEGG_ADHERENS_JUNCTION
GO_ENDOLYSOSOME, GO_ENDOLYSOSOME
GCNP_SHH_UP_EARLY.V1_DN, GCNP_SHH_UP_EARLY.V1_DN
BIOCARTA_BARRESTIN_SRC_PATHWAY, BIOCARTA_BARRESTIN_SRC_PATHWAY
MOOTHA_FFA_OXYDATION, MOOTHA_FFA_OXYDATION
BAUS_TFF2_TARGETS_UP, BAUS_TFF2_TARGETS_UP
GO_PEPTIDE_CATABOLIC_PROCESS, GO_PEPTIDE_CATABOLIC_PROCESS
GO_ACTIN_BASED_CELL_PROJECTION, GO_ACTIN_BASED_CELL_PROJECTION
BIOCARTA_BARR_MAPK_PATHWAY, BIOCARTA_BARR_MAPK_PATHWAY
GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP, GSE3720_LPS_VS_PMA_STIM_VD2_GAMMADELTA_TCELL_UP
GRANDVAUX_IFN_RESPONSE_NOT_VIA_IRF3, GRANDVAUX_IFN_RESPONSE_NOT_VIA_IRF3
GO_CELLULAR_RESPONSE_TO_OXYGEN_LEVELS, GO_CELLULAR_RESPONSE_TO_OXYGEN_LEVELS
GO_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS, GO_MEMBRANE_PROTEIN_ECTODOMAIN_PROTEOLYSIS
PID_ARF_3PATHWAY, PID_ARF_3PATHWAY
GO_ENDOLYSOSOME_MEMBRANE, GO_ENDOLYSOSOME_MEMBRANE
GO_CYTOPLASMIC_UBIQUITIN_LIGASE_COMPLEX, GO_CYTOPLASMIC_UBIQUITIN_LIGASE_COMPLEX
GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP, GSE40274_FOXP3_VS_FOXP3_AND_HELIOS_TRANSDUCED_ACTIVATED_CD4_TCELL_UP
PID_NCADHERIN_PATHWAY, PID_NCADHERIN_PATHWAY
MODULE_274, MODULE_274
GO_METALLOAMINOPEPTIDASE_ACTIVITY, GO_METALLOAMINOPEPTIDASE_ACTIVITY
GO_HINDBRAIN_MORPHOGENESIS, GO_HINDBRAIN_MORPHOGENESIS
GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_I_PROMOTER, GO_POSITIVE_REGULATION_OF_TRANSCRIPTION_FROM_RNA_POLYMERASE_I_PROMOTER
GO_CLATHRIN_BINDING, GO_CLATHRIN_BINDING
REACTOME_GAP_JUNCTION_DEGRADATION, REACTOME_GAP_JUNCTION_DEGRADATION