

S_16H_IFNG_IN_CD8POS_DC_DN, GSE3337_CTRL_VS_16H_IFNG_IN_CD8POS_DC_DN

- GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_DN, GSE46242_CTRL_VS_EGR2_DELETED_ANERGIC_TH1_CD4_TCELL_DN
- GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT, GO_NEGATIVE_REGULATION_OF_VASCULATURE_DEVELOPMENT
- GO_CHEMOKINE_MEDIATED_SIGNALING_PATHWAY, GO_CHEMOKINE_MEDIATED_SIGNALING_PATHWAY
- GO_CELLULAR_DEFENSE_RESPONSE, GO_CELLULAR_DEFENSE_RESPONSE
- HOFFMANN_LARGE_TO_SMALL_PRE_BII_LYMPHOCYTE_DN, HOFFMANN_LARGE_TO_SMALL_PRE_BII_LYMPHOCYTE_DN
- GO_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS, GO_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS
- CROMER_TUMORIGENESIS_UP, CROMER_TUMORIGENESIS_UP
- LEE_LIVER_CANCER_ACOX1_DN, LEE_LIVER_CANCER_ACOX1_DN
- GO_POSITIVE_REGULATION_OF_CAMP_MEDIATED_SIGNALING, GO_POSITIVE_REGULATION_OF_CAMP_MEDIATED_SIGNALING
- GO_POSITIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_NUCLEOTIDE_METABOLIC_PROCESS
- GO_POSITIVE_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS, GO_POSITIVE_REGULATION_OF_CYCLIC_NUCLEOTIDE_METABOLIC_PROCESS
- GO_REGULATION_OF_CAMP_METABOLIC_PROCESS, GO_REGULATION_OF_CAMP_METABOLIC_PROCESS
- GO_CAMERA_TYPE_EYE_PHOTORECEPTOR_CELL_DIFFERENTIATION, GO_CAMERA_TYPE_EYE_PHOTORECEPTOR_CELL_DIFFERENTIATION
- GO_COLLAGEN_FIBRIL_ORGANIZATION, GO_COLLAGEN_FIBRIL_ORGANIZATION