AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_DN, GSE2770\_IL12\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_DN

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GSE3982 CTRL VS LPS 4H MAC DN, GSE3982 CTRL VS LPS 4H MAC DN
HALLMARK INTERFERON ALPHA RESPONSE, HALLMARK INTERFERON ALPHA RESPONSE
TNCATNTCCYR_UNKNOWN, TNCATNTCCYR_UNKNOWN
REACTOME SIGNALING BY ILS, REACTOME SIGNALING BY ILS
GNF2 TPT1, GNF2 TPT1
GSE2706 2H_VS_8H_LPS_STIM_DC_DN, GSE2706_2H_VS_8H_LPS_STIM_DC_DN
AZARE_NEOPLASTIC_TRANSFORMATION_BY_STAT3_UP, AZARE_NEOPLASTIC_TRANSFORMATION_BY_STAT3_UP
GSE42021_CD24HI_VS_CD24LOW_TREG_THYMUS_DN, GSE42021_CD24HI_VS_CD24LOW_TREG_THYMUS_DN
GO_STEM_CELL_DIFFERENTIATION, GO_STEM_CELL_DIFFERENTIATION
PID CERAMIDE PATHWAY, PID CERAMIDE PATHWAY
KIM_MYC_AMPLIFICATION_TARGETS_DN, KIM_MYC_AMPLIFICATION_TARGETS_DN
GSE42021 TREG_PLN_VS_CD24LO_TREG_THYMUS_DN, GSE42021 TREG_PLN_VS_CD24LO_TREG_THYMUS_DN
SA B CELL RECEPTOR COMPLEXES, SA B CELL RECEPTOR COMPLEXES
GO ADENYLYLTRANSFERASE ACTIVITY, GO ADENYLYLTRANSFERASE ACTIVITY
BIOCARTA_CERAMIDE_PATHWAY, BIOCARTA_CERAMIDE_PATHWAY
GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN, GSE28783_ANTI_MIR33_VS_UNTREATED_ATHEROSCLEROSIS_MACROPHAGE_DN
GALLUZZI_PREVENT_MITOCHONDIAL_PERMEABILIZATION, GALLUZZI_PREVENT_MITOCHONDIAL_PERMEABILIZATION
GO_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY, GO_PHOSPHATIDYLINOSITOL_3_KINASE_ACTIVITY
PID_P38_ALPHA_BETA_PATHWAY, PID_P38_ALPHA_BETA_PATHWAY
GO_CELLULAR_RESPONSE_TO_MECHANICAL_STIMULUS, GO_CELLULAR_RESPONSE_TO_MECHANICAL_STIMULUS
SEITZ_NEOPLASTIC_TRANSFORMATION_BY_8P_DELETION_UP, SEITZ_NEOPLASTIC_TRANSFORMATION_BY_8P_DELETION_UP
MOSERLE_IFNA_RESPONSE, MOSERLE_IFNA_RESPONSE
GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_TRASDUCED_CD4_TCELL_DN, GSE37533_UNTREATED_VS_PIOGLIZATONE_TREATED_CD4_TCELL_FOXP3_T
REACTOME G PROTEIN BETA GAMMA SIGNALLING, REACTOME G PROTEIN BETA GAMMA SIGNALLING
GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_180MIN_UP, GSE5589_LPS_VS_LPS_AND_IL10_STIM_IL10_KO_MACROPHAGE_180MIN_UP
GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_UP, GSE22589_SIV_VS_HIV_AND_SIV_INFECTED_DC_UP
GO_POSITIVE_REGULATION_OF_LIPASE_ACTIVITY, GO_POSITIVE_REGULATION_OF_LIPASE_ACTIVITY
GO MUSCLE HYPERTROPHY, GO MUSCLE HYPERTROPHY
GSE22025_UNTREATED_VS_PROGESTERONE_TREATED_CD4_TCELL_UP, GSE22025_UNTREATED_VS_PROGESTERONE_TREATED_CD4_TCELL_UP
GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_DN, GSE19888_CTRL_VS_T_CELL_MEMBRANES_ACT_MAST_CELL_DN
GSE42088 2H VS 24H LEISHMANIA INF DC DN, GSE42088 2H VS 24H LEISHMANIA INF DC DN
GNCF 01, GNCF 01
ROETH_TERT_TARGETS_UP, ROETH_TERT_TARGETS_UP
ZHAN MULTIPLE MYELOMA HP_UP, ZHAN MULTIPLE MYELOMA HP_UP
GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION OF ESTABLISHMENT OF PROTEIN LOCALIZATION TO PLASMA MEMBRANE, GO NEGATIVE REGULATION TO PLASMA MEMBRANE, GO NEGATIV
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GSE21360\_NAIVE\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_UP, GSE21360\_NAIVE\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_UP