GSE17721\_LPS\_VS\_CPG\_8H\_BMDC\_DN, GSE17721\_LPS\_VS\_CPG\_8H\_BMDC\_DN GSE17721\_0.5H\_VS\_24H\_GARDIQUIMOD\_BMDC\_DN, GSE17721\_0.5H\_VS\_24H\_GARDIQUIMOD\_BMDC\_DN GSE17721\_POLYIC\_VS\_PAM3CSK4\_12H\_BMDC\_DN, GSE17721\_POLYIC\_VS\_PAM3CSK4\_12H\_BMDC\_DN GSE12198\_CTRL\_VS\_LOW\_IL2\_STIM\_NK\_CELL\_DN, GSE12198\_CTRL\_VS\_LOW\_IL2\_STIM\_NK\_CELL\_DN MORF\_RFC1, MORF\_RFC1 GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_12H\_BMDC\_UP, GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_12H\_BMDC\_UP GSE17721\_0.5H\_VS\_4H\_GARDIQUIMOD\_BMDC\_UP, GSE17721\_0.5H\_VS\_4H\_GARDIQUIMOD\_BMDC\_UP GSE17721\_CPG\_VS\_GARDIQUIMOD\_12H\_BMDC\_DN, GSE17721\_CPG\_VS\_GARDIQUIMOD\_12H\_BMDC\_DN GSE17721\_0.5H\_VS\_12H\_PAM3CSK4\_BMDC\_DN, GSE17721\_0.5H\_VS\_12H\_PAM3CSK4\_BMDC\_DN RB\_P107\_DN.V1\_DN, RB\_P107\_DN.V1\_DN GSE7852\_TREG\_VS\_TCONV\_UP, GSE7852\_TREG\_VS\_TCONV\_UP GSE33425\_CD161\_HIGH\_VS\_NEG\_CD8\_TCELL\_DN, GSE33425\_CD161\_HIGH\_VS\_NEG\_CD8\_TCELL\_DN GSE17721\_LPS\_VS\_PAM3CSK4\_0.5H\_BMDC\_UP, GSE17721\_LPS\_VS\_PAM3CSK4\_0.5H\_BMDC\_UP GSE14415\_INDUCED\_TREG\_VS\_FOXP3\_KO\_INDUCED\_TREG\_DN, GSE14415\_INDUCED\_TREG\_VS\_FOXP3\_KO\_INDUCED\_TREG\_DN GSE17721\_PAM3CSK4\_VS\_CPG\_1H\_BMDC\_DN, GSE17721\_PAM3CSK4\_VS\_CPG\_1H\_BMDC\_DN PID\_HEDGEHOG\_GLI\_PATHWAY, PID\_HEDGEHOG\_GLI\_PATHWAY GO PROTEIN COMPLEX SCAFFOLD, GO PROTEIN COMPLEX SCAFFOLD M\_STIM\_CD4\_TCELL\_ATOPIC\_PATIENT\_UP, GSE14908\_RESTING\_VS\_HDM\_STIM\_CD4\_TCELL\_ATOPIC\_PATIENT\_UP GO\_CARBON\_OXYGEN\_LYASE\_ACTIVITY, GO\_CARBON\_OXYGEN\_LYASE\_ACTIVITY GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_16H\_BMDC\_UP, GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_16H\_BMDC\_UP GO\_NEGATIVE\_REGULATION\_OF\_DEPHOSPHORYLATION, GO\_NEGATIVE\_REGULATION\_OF\_DEPHOSPHORYLATION NIKOLSKY\_BREAST\_CANCER\_16Q24\_AMPLICON, NIKOLSKY\_BREAST\_CANCER\_16Q24\_AMPLICON VANTVEER\_BREAST\_CANCER\_BRCA1\_UP, VANTVEER\_BREAST\_CANCER\_BRCA1\_UP CHEN\_LIVER\_METABOLISM\_QTL\_CIS, CHEN\_LIVER\_METABOLISM\_QTL\_CIS GO\_EXOPEPTIDASE\_ACTIVITY, GO\_EXOPEPTIDASE\_ACTIVITY GO\_SARCOPLASM, GO\_SARCOPLASM GO\_CILIARY\_TRANSITION\_ZONE, GO\_CILIARY\_TRANSITION\_ZONE GSE24972 WT VS IRF8 KO SPLEEN FOLLICULAR BCELL UP, GSE24972 WT VS IRF8 KO SPLEEN FOLLICULAR BCELL UP CAFFAREL\_RESPONSE\_TO\_THC\_24HR\_3\_UP, CAFFAREL\_RESPONSE\_TO\_THC\_24HR\_3\_UP GSE13887\_RESTING\_VS\_NO\_TREATED\_CD4\_TCELL\_DN, GSE13887\_RESTING\_VS\_NO\_TREATED\_CD4\_TCELL\_DN GO\_PROTEIN\_TARGETING\_TO\_LYSOSOME, GO\_PROTEIN\_TARGETING\_TO\_LYSOSOME GO\_ENDOPLASMIC\_RETICULUM\_SUBCOMPARTMENT, GO\_ENDOPLASMIC\_RETICULUM\_SUBCOMPARTMENT chr10p12, chr10p12 GSE13762\_CTRL\_VS\_125\_VITAMIND\_DAY12\_DC\_DN, GSE13762\_CTRL\_VS\_125\_VITAMIND\_DAY12\_DC\_DN GNF2\_MMP1, GNF2\_MMP1 GO\_PROTEIN\_LOCALIZATION\_TO\_LYSOSOME, GO\_PROTEIN\_LOCALIZATION\_TO\_LYSOSOME

GSE14908\_ATOPIC\_VS\_NONATOPIC\_PATIENT\_RESTING\_CD4\_TCELL\_DN, GSE14908\_ATOPIC\_VS\_NONATOPIC\_PATIENT\_RESTING\_CD4\_TCELL\_DN GSE23505\_UNTREATED\_VS\_4DAY\_IL6\_IL1\_TREATED\_CD4\_TCELL\_UP, GSE23505\_UNTREATED\_VS\_4DAY\_IL6\_IL1\_TREATED\_CD4\_TCELL\_UP GSE14908\_ATOPIC\_VS\_NONATOPIC\_PATIENT\_HDM\_STIM\_CD4\_TCELL\_UP, GSE14908\_ATOPIC\_VS\_NONATOPIC\_PATIENT\_HDM\_STIM\_CD4\_TCELL\_UP GSE15330\_LYMPHOID\_MULTIPOTENT\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_DN, GSE15330\_LYMPHOID\_MULTIPOTENT\_VS\_GRANULOCYTE\_MONOCYTE\_PRO GSE15330\_WT\_VS\_IKAROS\_KO\_MEGAKARYOCYTE\_ERYTHROID\_PROGENITOR\_UP, GSE15330\_WT\_VS\_IKAROS\_KO\_MEGAKARYOCYTE\_ERYTHROID\_PROGENITOR\_UP GSE9509\_10MIN\_VS\_30MIN\_LPS\_AND\_IL10\_STIM\_IL10\_KO\_MACROPHAGE\_DN, GSE9509\_10MIN\_VS\_30MIN\_LPS\_AND\_IL10\_STIM\_IL10\_KO\_MACROPHAGE\_DN GSE41978\_WT\_VS\_BIM\_KO\_KLRG1\_LOW\_EFFECTOR\_CD8\_TCELL\_UP, GSE41978\_WT\_VS\_BIM\_KO\_KLRG1\_LOW\_EFFECTOR\_CD8\_TCELL\_UP GO\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_VACUOLE, GO\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_VACUOLE GO\_POSITIVE\_REGULATION\_OF\_VIRAL\_RELEASE\_FROM\_HOST\_CELL, GO\_POSITIVE\_REGULATION\_OF\_VIRAL\_RELEASE\_FROM\_HOST\_CELL GO\_REGULATION\_OF\_EXTRACELLULAR\_MATRIX\_ORGANIZATION, GO\_REGULATION\_OF\_EXTRACELLULAR\_MATRIX\_ORGANIZATION GO\_NEGATIVE\_REGULATION\_OF\_PHOSPHOPROTEIN\_PHOSPHATASE\_ACTIVITY, GO\_NEGATIVE\_REGULATION\_OF\_PHOSPHOPROTEIN\_PHOSPHATASE\_ACTIVITY GO\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT\_PEPTIDE\_BONDS, GO\_HYDROLASE\_ACTIVITY\_ACTING\_ON\_CARBON\_NITROGEN\_BUT\_NOT