GOMF\_PROTEIN\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_PROTEIN\_METHYLTRANSFERASE\_ACTIVITY GSE15659\_NAIVE\_CD4\_TCELL\_VS\_RESTING\_TREG\_DN, GSE15659\_NAIVE\_CD4\_TCELL\_VS\_RESTING\_TREG\_DN NIKOLSKY\_BREAST\_CANCER\_16P13\_AMPLICON, NIKOLSKY\_BREAST\_CANCER\_16P13\_AMPLICON GSE16266\_CTRL\_VS\_LPS\_STIM\_MEF\_DN, GSE16266\_CTRL\_VS\_LPS\_STIM\_MEF\_DN GOBP\_PROTEIN\_ADP\_RIBOSYLATION, GOBP\_PROTEIN\_ADP\_RIBOSYLATION MORF\_PDPK1, MORF\_PDPK1 GSE37301\_LYMPHOID\_PRIMED\_MPP\_VS\_PRO\_BCELL\_DN, GSE37301\_LYMPHOID\_PRIMED\_MPP\_VS\_PRO\_BCELL\_DN GAUSSMANN\_MLL\_AF4\_FUSION\_TARGETS\_A\_DN, GAUSSMANN\_MLL\_AF4\_FUSION\_TARGETS\_A\_DN MIR4433B\_5P, MIR4433B\_5P GOMF\_DEACETYLASE\_ACTIVITY, GOMF\_DEACETYLASE\_ACTIVITY GNF2\_LCAT, GNF2\_LCAT GNF2\_HPN, GNF2\_HPN MIR6848\_5P, MIR6848\_5P OID VS MAC IL25 TREATED LUNG DN, GSE36392 TYPE 2 MYELOID VS MAC IL25 TREATED LUNG DN GNF2 GSTM1, GNF2 GSTM1 MIR3672, MIR3672 GOBP\_SNO\_S\_RNA\_3\_END\_PROCESSING, GOBP\_SNO\_S\_RNA\_3\_END\_PROCESSING GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_BINDING, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_BINDING GSE35543\_IN\_VIVO\_NTREG\_VS\_IN\_VITRO\_ITREG\_UP, GSE35543\_IN\_VIVO\_NTREG\_VS\_IN\_VITRO\_ITREG\_UP KAAB\_FAILED\_HEART\_ATRIUM\_UP, KAAB\_FAILED\_HEART\_ATRIUM\_UP EGR3\_01, EGR3\_01 GOBP\_HISTONE\_H4\_DEACETYLATION, GOBP\_HISTONE\_H4\_DEACETYLATION MIR6864\_3P, MIR6864\_3P GNF2\_TST, GNF2\_TST GOBP\_REGULATION\_OF\_LIPASE\_ACTIVITY, GOBP\_REGULATION\_OF\_LIPASE\_ACTIVITY HP HYDROXYPROLINURIA, HP HYDROXYPROLINURIA MIR6890 5P, MIR6890 5P FOXC1 TARGET GENES, FOXC1 TARGET GENES JOHANSSON\_BRAIN\_CANCER\_EARLY\_VS\_LATE\_DN, JOHANSSON\_BRAIN\_CANCER\_EARLY\_VS\_LATE\_DN HP\_NEPHRONOPHTHISIS, HP\_NEPHRONOPHTHISIS MODULE\_458, MODULE\_458 HP\_ABNORMAL\_RENAL\_MEDULLA\_MORPHOLOGY, HP\_ABNORMAL\_RENAL\_MEDULLA\_MORPHOLOGY GOCC\_AXONAL\_GROWTH\_CONE, GOCC\_AXONAL\_GROWTH\_CONE GOBP\_DETOXIFICATION\_OF\_NITROGEN\_COMPOUND, GOBP\_DETOXIFICATION\_OF\_NITROGEN\_COMPOUND GOBP\_POLYSACCHARIDE\_CATABOLIC\_PROCESS, GOBP\_POLYSACCHARIDE\_CATABOLIC\_PROCESS MIZUKAMI\_HYPOXIA\_DN, MIZUKAMI\_HYPOXIA\_DN GOBP\_NEGATIVE\_REGULATION\_OF\_MUSCLE\_ORGAN\_DEVELOPMENT, GOBP\_NEGATIVE\_REGULATION\_OF\_MUSCLE\_ORGAN\_DEVELOPMENT

GSE16386 IL4 VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_MACROPHAGE\_6H\_UP, GSE16386\_IL4\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_MACROPHAGE\_6H\_UP GSE36392\_TYPE\_2\_MYELOID\_VS\_NEUTROPHIL\_IL25\_TREATED\_LUNG\_DN, GSE36392\_TYPE\_2\_MYELOID\_VS\_NEUTROPHIL\_IL25\_TREATED\_LUNG\_DN GSE37533\_UNTREATED\_VS\_PIOGLIZATONE\_TREATED\_CD4\_TCELL\_PPARG2\_AND\_FOXP3\_TRASDUCED\_DN, GSE37533\_UNTREATED\_VS\_PIOGLIZATONE\_TREATED\_CD4\_TCELL\_PPARG2\_AND\_FOXP3\_TRASDUCED\_DN GSE3039\_NKT\_CELL\_VS\_ALPHABETA\_CD8\_TCELL\_DN, GSE3039\_NKT\_CELL\_VS\_ALPHABETA\_CD8\_TCELL\_DN GSE25123\_ROSIGLITAZONE\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_PPARG\_KO\_MACROPHAGE\_DAY10\_DN, GSE25123\_ROSIGLITAZONE\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_PPARG\_KO\_MACROPHAGE\_DAY10\_DN GSE45365\_HEALTHY\_VS\_MCMV\_INFECTION\_CD11B\_DC\_IFNAR\_KO\_UP, GSE45365\_HEALTHY\_VS\_MCMV\_INFECTION\_CD11B\_DC\_IFNAR\_KO\_UP GSE10240\_IL17\_VS\_IL17\_AND\_IL22\_STIM\_PRIMARY\_BRONCHIAL\_EPITHELIAL\_CELLS\_DN, GSE10240\_IL17\_VS\_IL17\_AND\_IL22\_STIM\_PRIMARY\_BRONCHIAL\_EPITHELIAL\_CELLS\_DN GSE9509\_10MIN\_VS\_30MIN\_LPS\_STIM\_IL10\_KO\_MACROPHAGE\_DN, GSE9509\_10MIN\_VS\_30MIN\_LPS\_STIM\_IL10\_KO\_MACROPHAGE\_DN GSE15659 CD45RA\_NEG\_CD4\_TCELL\_VS\_RESTING\_TREG\_DN, GSE15659\_CD45RA\_NEG\_CD4\_TCELL\_VS\_RESTING\_TREG\_DN GSE19888\_ADENOSINE\_A3R\_INH\_VS\_ACT\_WITH\_INHIBITOR\_PRETREATMENT\_IN\_MAST\_CELL\_DN, GSE19888\_ADENOSINE\_A3R\_INH\_VS\_ACT\_WITH\_INHIBITOR\_PRETREATMENT\_IN\_MAST\_CELL\_DN GSE9650\_GP33\_VS\_GP276\_LCMV\_SPECIFIC\_EXHAUSTED\_CD8\_TCELL\_UP, GSE9650\_GP33\_VS\_GP276\_LCMV\_SPECIFIC\_EXHAUSTED\_CD8\_TCELL\_UP GOMF\_NADPLUS\_ADP\_RIBOSYLTRANSFERASE\_ACTIVITY, GOMF\_NADPLUS\_ADP\_RIBOSYLTRANSFERASE\_ACTIVITY HOEK\_NK\_CELL\_2011\_2012\_TIV\_1D\_VS\_0DY\_ADULT\_1D\_DN, HOEK\_NK\_CELL\_2011\_2012\_TIV\_1D\_VS\_0DY\_ADULT\_1D\_DN GSE22935\_UNSTIM\_VS\_48H\_MBOVIS\_BCG\_STIM\_MYD88\_KO\_MACROPHAGE\_DN, GSE22935\_UNSTIM\_VS\_48H\_MBOVIS\_BCG\_STIM\_MYD88\_KO\_MACROPHAGE\_DN GAO\_LARGE\_INTESTINE\_24W\_C6\_SECRETORY\_PROGENITOR, GAO\_LARGE\_INTESTINE\_24W\_C6\_SECRETORY\_PROGENITOR GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_ACETYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_ACETYLATION FRANCO\_BLOOD\_SANOFI\_PASTEUR\_SA\_INACTIVATED\_INFLUENZA\_VACCINE\_CORRELATED\_WITH\_ANTIBODY\_RESPONSE\_AGE\_18\_40YO\_0DY\_POSITIVE, FRANCO\_BLOOD\_SANOFI\_PASTEUR\_SA\_INACTIVATED\_INFLUENZA\_VACCINE\_CORRELATED\_WITH\_ANTIBODY\_POSITIVE, FRANCO\_BLOOD\_SANOFI\_PASTEUR\_SA\_INACTIVATED\_INFLUENZA\_VACCINE\_CORRELATED\_WITH\_ANTIBODY\_POSITIVE, FRANCO\_BLOOD\_SANOFI\_PASTEUR\_SA\_INACTIVATED\_INFLUENZA\_VACCINE\_CORRELATED\_WITH\_ANTIBODY\_POSITIVE, FRANCO\_BLOOD\_POSITIVE, FRANCO\_ GOBP\_NEGATIVE\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_MEDIATED\_SIGNALING\_PATHWAY, GOBP\_NEGATIVE\_REGULATION\_OF\_TYPE\_I\_INTERFERON\_MEDIATED\_SIGNALING\_PATHWAY WP\_THE\_OVERLAP\_BETWEEN\_SIGNAL\_TRANSDUCTION\_PATHWAYS\_THAT\_CONTRIBUTE\_TO\_A\_RANGE\_OF\_LMNA\_LAMINOPATHIES, WP\_THE\_OVERLAP\_BETWEEN\_SIGNAL\_TRANSDUCTION\_PATHWAYS\_THAT\_CONTRIBUTE\_TO\_A\_RANGE\_OF\_LMNA\_LAMINOPATHIES GOBP\_CHEMOKINE\_C\_X\_C\_MOTIF\_LIGAND\_2\_PRODUCTION, GOBP\_CHEMOKINE\_C\_X\_C\_MOTIF\_LIGAND\_2\_PRODUCTION