GSE6674\_CPG\_VS\_PL2\_3\_STIM\_BCELL\_UP, GSE6674\_CPG\_VS\_PL2\_3\_STIM\_BCELL\_UP GSE17721\_LPS\_VS\_CPG\_4H\_BMDC\_DN, GSE17721\_LPS\_VS\_CPG\_4H\_BMDC\_DN GSE31082 DN\_VS\_CD4\_SP\_THYMOCYTE\_DN, GSE31082\_DN\_VS\_CD4\_SP\_THYMOCYTE\_DN GSE18893\_TCONV\_VS\_TREG\_24H\_TNF\_STIM\_DN, GSE18893\_TCONV\_VS\_TREG\_24H\_TNF\_STIM\_DN GSE17721\_CTRL\_VS\_PAM3CSK4\_6H\_BMDC\_UP, GSE17721\_CTRL\_VS\_PAM3CSK4\_6H\_BMDC\_UP GSE17721\_ALL\_VS\_24H\_PAM3CSK4\_BMDC\_DN, GSE17721\_ALL\_VS\_24H\_PAM3CSK4\_BMDC\_DN GSE20715\_0H\_VS\_24H\_OZONE\_LUNG\_UP, GSE20715\_0H\_VS\_24H\_OZONE\_LUNG\_UP GSE42021\_CD24HI\_VS\_CD24INT\_TREG\_THYMUS\_UP, GSE42021\_CD24HI\_VS\_CD24INT\_TREG\_THYMUS\_UP GSE13306\_TREG\_VS\_TCONV\_LAMINA\_PROPRIA\_UP, GSE13306\_TREG\_VS\_TCONV\_LAMINA\_PROPRIA\_UP GSE12198 CTRL VS HIGH IL2 STIM NK CELL DN, GSE12198 CTRL VS HIGH IL2 STIM NK CELL DN GSE18893\_CTRL\_VS\_TNF\_TREATED\_TCONV\_24H\_DN, GSE18893\_CTRL\_VS\_TNF\_TREATED\_TCONV\_24H\_DN GSE32128\_INOS\_DEPENDENT\_VS\_INOS\_INDEPENDENT\_ACTIVATED\_TCELL\_DN, GSE32128\_INOS\_DEPENDENT\_VS\_INOS\_INDEPENDENT\_ACTIVATED\_TCELL\_DN MIR124 5P, MIR124 5P TRAVAGLINI\_LUNG\_NATURAL\_KILLER\_CELL, TRAVAGLINI\_LUNG\_NATURAL\_KILLER\_CELL GSE26928\_EFF\_MEM\_VS\_CENTR\_MEM\_CD4\_TCELL\_UP, GSE26928\_EFF\_MEM\_VS\_CENTR\_MEM\_CD4\_TCELL\_UP GOBP\_PROTEIN\_LOCALIZATION\_TO\_VACUOLE, GOBP\_PROTEIN\_LOCALIZATION\_TO\_VACUOLE REACTOME\_ENDOSOMAL\_SORTING\_COMPLEX\_REQUIRED\_FOR\_TRANSPORT\_ESCRT, REACTOME\_ENDOSOMAL\_SORTING\_COMPLEX\_REQUIRED\_FOR\_TRANSPORT\_ESCRT MIR8089, MIR8089 ROSS AML WITH AML1 ETO FUSION, ROSS AML WITH AML1 ETO FUSION GOBP\_MULTIVESICULAR\_BODY\_ORGANIZATION, GOBP\_MULTIVESICULAR\_BODY\_ORGANIZATION GOCC\_ESCRT\_COMPLEX, GOCC\_ESCRT\_COMPLEX GOBP\_ENDOSOME\_TRANSPORT\_VIA\_MULTIVESICULAR\_BODY\_SORTING\_PATHWAY, GOBP\_ENDOSOME\_TRANSPORT\_VIA\_MULTIVESICULAR\_BODY\_SORTING\_PATHWAY MIR664A\_5P, MIR664A\_5P HP\_DEMYELINATING\_PERIPHERAL\_NEUROPATHY, HP\_DEMYELINATING\_PERIPHERAL\_NEUROPATHY ACT\_VS\_ACT\_CD4\_TCELL\_2H\_UP, GSE2770\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_2H\_UP BOYLAN\_MULTIPLE\_MYELOMA\_D\_CLUSTER\_DN, BOYLAN\_MULTIPLE\_MYELOMA\_D\_CLUSTER\_DN GOBP\_MULTIVESICULAR\_BODY\_SORTING\_PATHWAY, GOBP\_MULTIVESICULAR\_BODY\_SORTING\_PATHWAY MIR637, MIR637 HOFT CD4 POSITIVE ALPHA BETA MEMORY T CELL BCG VACCINE AGE 18 45YO ID 56D TOP 100 DEG AFTER IN VITRO RE STIMULATION UP, HOFT CD4 POSITIVE ALPHA BETA MEMORY T CELL BCG VACCINE AGE 18 45YO ID 56D TOP 100 DEG AFTER IN VITRO RE GOBP\_RESPIRATORY\_SYSTEM\_PROCESS, GOBP\_RESPIRATORY\_SYSTEM\_PROCESS ZNF502\_TARGET\_GENES, ZNF502\_TARGET\_GENES HP\_TYPE\_II\_LISSENCEPHALY, HP\_TYPE\_II\_LISSENCEPHALY HP\_CROHN\_S\_DISEASE, HP\_CROHN\_S\_DISEASE KEGG\_PRIMARY\_BILE\_ACID\_BIOSYNTHESIS, KEGG\_PRIMARY\_BILE\_ACID\_BIOSYNTHESIS GOBP\_RESPONSE\_TO\_REDOX\_STATE, GOBP\_RESPONSE\_TO\_REDOX\_STATE GOBP\_NEGATIVE\_REGULATION\_OF\_MULTI\_ORGANISM\_PROCESS, GOBP\_NEGATIVE\_REGULATION\_OF\_MULTI\_ORGANISM\_PROCESS GOBP\_OLIGOSACCHARIDE\_CATABOLIC\_PROCESS, GOBP\_OLIGOSACCHARIDE\_CATABOLIC\_PROCESS GOBP\_PIGMENT\_GRANULE\_ORGANIZATION, GOBP\_PIGMENT\_GRANULE\_ORGANIZATION HP\_TRACHEAL\_STENOSIS, HP\_TRACHEAL\_STENOSIS MIR6802\_5P, MIR6802\_5P MAGRANGEAS\_MULTIPLE\_MYELOMA\_IGLL\_VS\_IGLK\_DN, MAGRANGEAS\_MULTIPLE\_MYELOMA\_IGLL\_VS\_IGLK\_DN chr4p14, chr4p14 GOBP\_MORPHOGENESIS\_OF\_AN\_ENDOTHELIUM, GOBP\_MORPHOGENESIS\_OF\_AN\_ENDOTHELIUM DESCARTES\_MAIN\_FETAL\_NEUROENDOCRINE\_CELLS, DESCARTES\_MAIN\_FETAL\_NEUROENDOCRINE\_CELLS GOBP\_POSITIVE\_REGULATION\_OF\_MYOBLAST\_FUSION, GOBP\_POSITIVE\_REGULATION\_OF\_MYOBLAST\_FUSION GOBP\_REGULATION\_OF\_MYOBLAST\_FUSION, GOBP\_REGULATION\_OF\_MYOBLAST\_FUSION GOMF CADHERIN BINDING INVOLVED IN CELL CELL ADHESION, GOMF CADHERIN BINDING INVOLVED IN CELL CELL ADHESION REACTOME\_THE\_ACTIVATION\_OF\_ARYLSULFATASES, REACTOME\_THE\_ACTIVATION\_OF\_ARYLSULFATASES DESCARTES\_MAIN\_FETAL\_INHIBITORY\_NEURONS, DESCARTES\_MAIN\_FETAL\_INHIBITORY\_NEURONS