

GSE32986\_UNSTIM\_VS\_CURDLAN\_HIGHDOSE\_STIM\_DC\_DN, GSE32986\_UNSTIM\_VS\_CURDLAN\_HIGHDOSE\_STIM\_DC\_DN  
GSE32986\_UNSTIM\_VS\_CURDLAN\_LOWDOSE\_STIM\_DC\_DN, GSE32986\_UNSTIM\_VS\_CURDLAN\_LOWDOSE\_STIM\_DC\_DN  
GSE40068\_BCL6\_POS\_VS\_NEG\_CXCR5\_POS\_TFH\_DN, GSE40068\_BCL6\_POS\_VS\_NEG\_CXCR5\_POS\_TFH\_DN  
GSE26495\_NAIVE\_VS\_PD1LOW\_CD8\_TCELL\_DN, GSE26495\_NAIVE\_VS\_PD1LOW\_CD8\_TCELL\_DN  
GSE26495\_NAIVE\_VS\_PD1HIGH\_CD8\_TCELL\_DN, GSE26495\_NAIVE\_VS\_PD1HIGH\_CD8\_TCELL\_DN  
GSE24574\_BCL6\_HIGH\_VS\_LOW\_TFH\_CD4\_TCELL\_UP, GSE24574\_BCL6\_HIGH\_VS\_LOW\_TFH\_CD4\_TCELL\_UP  
GSE5099\_DAY3\_VS\_DAY7\_MCSF\_TREATED\_MACROPHAGE\_UP, GSE5099\_DAY3\_VS\_DAY7\_MCSF\_TREATED\_MACROPHAGE\_UP  
GSE30962\_ACUTE\_VS\_CHRONIC\_LCMV\_SECONDARY\_INF\_CD8\_TCELL\_UP, GSE30962\_ACUTE\_VS\_CHRONIC\_LCMV\_SECONDARY\_INF\_CD8\_TCELL\_UP  
GSE11057\_NAIVE\_VS\_MEMORY\_CD4\_TCELL\_DN, GSE11057\_NAIVE\_VS\_MEMORY\_CD4\_TCELL\_DN  
GSE6269\_HEALTHY\_VS\_STAPH\_PNEUMO\_INF\_PBMC\_UP, GSE6269\_HEALTHY\_VS\_STAPH\_PNEUMO\_INF\_PBMC\_UP  
GSE7460\_CD8\_TCELL\_VS\_CD4\_TCELL\_ACT\_UP, GSE7460\_CD8\_TCELL\_VS\_CD4\_TCELL\_ACT\_UP  
GSE32986\_UNSTIM\_VS\_GMCSF\_AND\_CURDLAN\_HIGHDOSE\_STIM\_DC\_DN, GSE32986\_UNSTIM\_VS\_GMCSF\_AND\_CURDLAN\_HIGHDOSE\_STIM\_DC\_DN  
GSE5542\_UNTREATED\_VS\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP, GSE5542\_UNTREATED\_VS\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP  
SREBP1\_01, SREBP1\_01  
AIZARANI\_LIVER\_C1\_NK\_NKT\_CELLS\_1, AIZARANI\_LIVER\_C1\_NK\_NKT\_CELLS\_1  
GSE40666\_UNTREATED\_VS\_IFNA\_STIM\_EFFECTOR\_CD8\_TCELL\_90MIN\_UP, GSE40666\_UNTREATED\_VS\_IFNA\_STIM\_EFFECTOR\_CD8\_TCELL\_90MIN\_UP  
GSE2935\_UV\_INACTIVATED\_VS\_LIVE\_SENDAI\_VIRUS\_INF\_MACROPHAGE\_DN, GSE2935\_UV\_INACTIVATED\_VS\_LIVE\_SENDAI\_VIRUS\_INF\_MACROPHAGE\_DN  
GSE26669\_CTRL\_VS\_COSTIM\_BLOCK\_MLR\_CD8\_TCELL\_DN, GSE26669\_CTRL\_VS\_COSTIM\_BLOCK\_MLR\_CD8\_TCELL\_DN  
GSE34205\_RSV\_VS\_FLU\_INF\_INFANT\_PBMC\_DN, GSE34205\_RSV\_VS\_FLU\_INF\_INFANT\_PBMC\_DN  
GSE11057\_NAIVE\_VS\_CENT\_MEMORY\_CD4\_TCELL\_DN, GSE11057\_NAIVE\_VS\_CENT\_MEMORY\_CD4\_TCELL\_DN  
GSE37416\_12H\_VS\_48H\_F\_TULARENSIS\_LV5\_NEUTROPHIL\_DN, GSE37416\_12H\_VS\_48H\_F\_TULARENSIS\_LV5\_NEUTROPHIL\_DN  
GSE13738\_RESTING\_VS\_BYSTANDER\_ACTIVATED\_CD4\_TCELL\_UP, GSE13738\_RESTING\_VS\_BYSTANDER\_ACTIVATED\_CD4\_TCELL\_UP  
GSE7852\_TREG\_VS\_TCONV\_THYMUS\_DN, GSE7852\_TREG\_VS\_TCONV\_THYMUS\_DN  
GSE21379\_WT\_VS\_SAP\_KO\_TFH\_CD4\_TCELL\_DN, GSE21379\_WT\_VS\_SAP\_KO\_TFH\_CD4\_TCELL\_DN  
GSE17721\_POLYIC\_VS\_PAM3CSK4\_6H\_BMDC\_UP, GSE17721\_POLYIC\_VS\_PAM3CSK4\_6H\_BMDC\_UP  
GSE44649\_WT\_VS\_MIR155\_KO\_NAIVE\_CD8\_TCELL\_UP, GSE44649\_WT\_VS\_MIR155\_KO\_NAIVE\_CD8\_TCELL\_UP  
GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_12H\_BMDC\_DN, GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_12H\_BMDC\_DN  
GSE3039\_NKT\_CELL\_VS\_ALPHAALPHA\_CD8\_TCELL\_DN, GSE3039\_NKT\_CELL\_VS\_ALPHAALPHA\_CD8\_TCELL\_DN  
GSE5542\_UNTREATED\_VS\_IFNA\_TREATED\_EPITHELIAL\_CELLS\_6H\_UP, GSE5542\_UNTREATED\_VS\_IFNA\_TREATED\_EPITHELIAL\_CELLS\_6H\_UP  
GSE13738\_RESTING\_VS\_TCR\_ACTIVATED\_CD4\_TCELL\_UP, GSE13738\_RESTING\_VS\_TCR\_ACTIVATED\_CD4\_TCELL\_UP  
GSE17721\_CPG\_VS\_GARDIQUIMOD\_0.5H\_BMDC\_DN, GSE17721\_CPG\_VS\_GARDIQUIMOD\_0.5H\_BMDC\_DN  
GSE3982\_CTRL\_VS\_IGE\_STIM\_MAST\_CELL\_UP, GSE3982\_CTRL\_VS\_IGE\_STIM\_MAST\_CELL\_UP  
GSE43863\_LY6C\_INT\_CXCR5POS\_VS\_LY6C\_LOW\_CXCR5NEG\_EFFECTOR\_CD4\_TCELL\_DN, GSE43863\_LY6C\_INT\_CXCR5POS\_VS\_LY6C\_LOW\_CXCR5NEG\_EFFECTOR\_CD4\_TCELL\_DN  
GSE17721\_12H\_VS\_24H\_GARDIQUIMOD\_BMDC\_UP, GSE17721\_12H\_VS\_24H\_GARDIQUIMOD\_BMDC\_UP  
GSE3720\_VD1\_VS\_VD2\_GAMMADELTA\_TCELL\_WITH\_LPS\_STIM\_UP, GSE3720\_VD1\_VS\_VD2\_GAMMADELTA\_TCELL\_WITH\_LPS\_STIM\_UP  
GSE1925\_CTRL\_VS\_24H\_IFNG\_STIM\_MACROPHAGE\_UP, GSE1925\_CTRL\_VS\_24H\_IFNG\_STIM\_MACROPHAGE\_UP  
GSE5455\_HEALTHY\_VS\_TUMOR\_BEARING\_MOUSE\_SPLEEN\_MONOCYTE\_UP, GSE5455\_HEALTHY\_VS\_TUMOR\_BEARING\_MOUSE\_SPLEEN\_MONOCYTE\_UP  
GSE32986\_GMCSF\_VS\_GMCSF\_AND\_CURDLAN\_HIGHDOSE\_STIM\_DC\_DN, GSE32986\_GMCSF\_VS\_GMCSF\_AND\_CURDLAN\_HIGHDOSE\_STIM\_DC\_DN  
HP\_HIRSUTISM, HP\_HIRSUTISM  
ODONNELL\_TARGETS\_OF\_MYC\_AND\_TFRC\_UP, ODONNELL\_TARGETS\_OF\_MYC\_AND\_TFRC\_UP  
GSE43955\_10H\_VS\_30H\_ACT\_CD4\_TCELL\_WITH\_TGFB\_IL6\_DN, GSE43955\_10H\_VS\_30H\_ACT\_CD4\_TCELL\_WITH\_TGFB\_IL6\_DN  
GSE22601\_DOUBLE\_NEGATIVE\_VS\_DOUBLE\_POSITIVE\_THYMOCYTE\_UP, GSE22601\_DOUBLE\_NEGATIVE\_VS\_DOUBLE\_POSITIVE\_THYMOCYTE\_UP  
HP\_HYDROPS\_FETALIS, HP\_HYDROPS\_FETALIS  
GSE35435\_RESTING\_VS\_IL4\_TREATED\_MACROPHAGE\_DN, GSE35435\_RESTING\_VS\_IL4\_TREATED\_MACROPHAGE\_DN  
GSE21033\_CTRL\_VS\_POLYIC\_STIM\_DC\_6H\_DN, GSE21033\_CTRL\_VS\_POLYIC\_STIM\_DC\_6H\_DN  
REACTOME\_GLYCOSAMINOGLYCAN\_METABOLISM, REACTOME\_GLYCOSAMINOGLYCAN\_METABOLISM  
STTTCRNTTT\_IRF\_Q6, STTTCRNTTT\_IRF\_Q6  
ZNF704\_TARGET\_GENES, ZNF704\_TARGET\_GENES  
GSE21033\_3H\_VS\_24H\_POLYIC\_STIM\_DC\_UP, GSE21033\_3H\_VS\_24H\_POLYIC\_STIM\_DC\_UP  
GSE16385\_UNTREATED\_VS\_12H\_IL4\_TREATED\_MACROPHAGE\_UP, GSE16385\_UNTREATED\_VS\_12H\_IL4\_TREATED\_MACROPHAGE\_UP  
GSE42088\_UNINF\_VS\_LEISHMANIA\_INF\_DC\_8H\_DN, GSE42088\_UNINF\_VS\_LEISHMANIA\_INF\_DC\_8H\_DN  
GSE3982\_EFF\_MEMORY\_CD4\_TCELL\_VS\_TH1\_UP, GSE3982\_EFF\_MEMORY\_CD4\_TCELL\_VS\_TH1\_UP  
REACTOME\_SPHINGOLIPID\_METABOLISM, REACTOME\_SPHINGOLIPID\_METABOLISM  
GSE3982\_EFF\_MEMORY\_CD4\_TCELL\_VS\_TH2\_UP, GSE3982\_EFF\_MEMORY\_CD4\_TCELL\_VS\_TH2\_UP  
GSE2585\_THYMIC\_MACROPHAGE\_VS\_MTEC\_UP, GSE2585\_THYMIC\_MACROPHAGE\_VS\_MTEC\_UP  
MIR7162\_5P, MIR7162\_5P  
REACTOME\_GLYCOSPHINGOLIPID\_METABOLISM, REACTOME\_GLYCOSPHINGOLIPID\_METABOLISM  
GOMF\_LIPASE\_ACTIVITY, GOMF\_LIPASE\_ACTIVITY  
HFH3\_01, HFH3\_01  
MIR6165, MIR6165  
GOBP\_REGULATION\_OF\_TORC1\_SIGNALING, GOBP\_REGULATION\_OF\_TORC1\_SIGNALING  
HP\_DYSMETRIA, HP\_DYSMETRIA  
BURTON\_ADIPOGENESIS\_9, BURTON\_ADIPOGENESIS\_9  
HP\_ARTERIAL\_STENOSIS, HP\_ARTERIAL\_STENOSIS  
GOBP\_AMINO\_ACID\_TRANSPORT, GOBP\_AMINO\_ACID\_TRANSPORT  
GSE3203\_HEALTHY\_VS\_INFLUENZA\_INFECTED\_LN\_BCELL\_DN, GSE3203\_HEALTHY\_VS\_INFLUENZA\_INFECTED\_LN\_BCELL\_DN  
GSE13547\_CTRL\_VS\_ANTLJGM\_STIM\_BCELL\_2H\_DN, GSE13547\_CTRL\_VS\_ANTLJGM\_STIM\_BCELL\_2H\_DN  
GSE360\_L\_MAJOR\_VS\_M\_TUBERCULOSIS\_MAC\_UP, GSE360\_L\_MAJOR\_VS\_M\_TUBERCULOSIS\_MAC\_UP  
GSE4142\_NAIVE\_VS\_MEMORY\_BCELL\_UP, GSE4142\_NAIVE\_VS\_MEMORY\_BCELL\_UP  
MODULE\_209, MODULE\_209  
GOBP\_POSITIVE\_REGULATION\_OF\_TOR\_SIGNALING, GOBP\_POSITIVE\_REGULATION\_OF\_TOR\_SIGNALING  
MIR6857\_5P, MIR6857\_5P  
HP\_ABNORMALITY\_OF\_OCULAR\_SMOOTH\_PURSUIT, HP\_ABNORMALITY\_OF\_OCULAR\_SMOOTH\_PURSUIT  
HP\_DIMINISHED\_MOTIVATION, HP\_DIMINISHED\_MOTIVATION  
GOBP\_CARBOHYDRATE\_DERIVATIVE\_TRANSPORT, GOBP\_CARBOHYDRATE\_DERIVATIVE\_TRANSPORT  
DESCARTES\_FETAL\_EYE\_SMOOTH\_MUSCLE\_CELLS, DESCARTES\_FETAL\_EYE\_SMOOTH\_MUSCLE\_CELLS  
HP\_ABNORMALITY\_OF\_THE\_VERTEBRAL\_ENDPLATES, HP\_ABNORMALITY\_OF\_THE\_VERTEBRAL\_ENDPLATES  
ZNF708\_TARGET\_GENES, ZNF708\_TARGET\_GENES  
GSE40274\_CTRL\_VS\_FOXP3\_AND\_SATB1\_TRANSDUCEDED\_ACTIVATED\_CD4\_TCELL\_DN, GSE40274\_CTRL\_VS\_FOXP3\_AND\_SATB1\_TRANSDUCEDED\_ACTIVATED\_CD4\_TCELL\_DN  
GSE5589\_WT\_VS\_IL10\_KO\_LPS\_STIM\_MACROPHAGE\_180MIN\_DN, GSE5589\_WT\_VS\_IL10\_KO\_LPS\_STIM\_MACROPHAGE\_180MIN\_DN  
ATOH8\_TARGET\_GENES, ATOH8\_TARGET\_GENES  
MIR4688, MIR4688  
REACTOME\_HEPARAN\_SULFATE\_HEPARIN\_HS\_GAG\_METABOLISM, REACTOME\_HEPARAN\_SULFATE\_HEPARIN\_HS\_GAG\_METABOLISM  
BIOCARTA\_CALCINEURIN\_PATHWAY, BIOCARTA\_CALCINEURIN\_PATHWAY  
REACTOME\_THE\_ACTIVATION\_OF\_ARYLSULFATASES, REACTOME\_THE\_ACTIVATION\_OF\_ARYLSULFATASES  
GOBP\_MORPHOGENESIS\_OF\_AN\_EPITHELIAL\_SHEET, GOBP\_MORPHOGENESIS\_OF\_AN\_EPITHELIAL\_SHEET  
HP\_PETECHIAE, HP\_PETECHIAE  
GOMF\_VOLUME\_SENSITIVE\_ANION\_CHANNEL\_ACTIVITY, GOMF\_VOLUME\_SENSITIVE\_ANION\_CHANNEL\_ACTIVITY  
MIR6743\_5P, MIR6743\_5P  
HP\_PROLONGED\_NEONATAL\_JAUNDICE, HP\_PROLONGED\_NEONATAL\_JAUNDICE  
GOBP\_AMINO\_SUGAR\_METABOLIC\_PROCESS, GOBP\_AMINO\_SUGAR\_METABOLIC\_PROCESS  
GOMF\_PROTEIN\_SELF\_ASSOCIATION, GOMF\_PROTEIN\_SELF\_ASSOCIATION  
HP\_DYSTROPHIC\_TOENAIL, HP\_DYSTROPHIC\_TOENAIL  
GOBP\_KERATAN\_SULFATE\_METABOLIC\_PROCESS, GOBP\_KERATAN\_SULFATE\_METABOLIC\_PROCESS  
GOMF\_ANION\_CHANNEL\_ACTIVITY, GOMF\_ANION\_CHANNEL\_ACTIVITY  
GOMF\_SULFURIC\_ESTER\_HYDROLASE\_ACTIVITY, GOMF\_SULFURIC\_ESTER\_HYDROLASE\_ACTIVITY  
FAN\_EMBRYONIC\_CTX\_EX\_2\_EXCITATORY\_NEURON, FAN\_EMBRYONIC\_CTX\_EX\_2\_EXCITATORY\_NEURON  
PETRETTO\_CARDIAC\_HYPERTROPHY, PETRETTO\_CARDIAC\_HYPERTROPHY  
MIR675\_3P, MIR675\_3P  
GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_GREY\_UP, GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_GREY\_UP  
GOBP\_ASSOCIATIVE\_LEARNING, GOBP\_ASSOCIATIVE\_LEARNING  
HP\_XANTHOMATOSIS, HP\_XANTHOMATOSIS  
GOBP\_POSITIVE\_REGULATION\_OF\_BONE\_MINERALIZATION, GOBP\_POSITIVE\_REGULATION\_OF\_BONE\_MINERALIZATION  
GOBP\_PHOSPHATIDYLETHANOLAMINE\_ACYL\_CHAIN\_REMODELING, GOBP\_PHOSPHATIDYLETHANOLAMINE\_ACYL\_CHAIN\_REMODELING  
HP\_ABNORMAL\_GERM\_CELL\_MORPHOLOGY, HP\_ABNORMAL\_GERM\_CELL\_MORPHOLOGY  
GOBP\_REGULATION\_OF\_STORE\_OPERATED\_CALCIIUM\_ENTRY, GOBP\_REGULATION\_OF\_STORE\_OPERATED\_CALCIIUM\_ENTRY  
WP\_GPR40\_PATHWAY, WP\_GPR40\_PATHWAY  
GOBP\_T\_HELPER\_17\_CELL\_LINEAGE\_COMMITMENT, GOBP\_T\_HELPER\_17\_CELL\_LINEAGE\_COMMITMENT  
MIR517A\_3P, MIR517B\_3P, MIR517C\_3P, MIR517A\_3P, MIR517B\_3P, MIR517C\_3P  
GOBP\_POSITIVE\_REGULATION\_OF\_OSTEOBLAST\_DIFFERENTIATION, GOBP\_POSITIVE\_REGULATION\_OF\_OSTEOBLAST\_DIFFERENTIATION  
REACTOME\_ACYL\_CHAIN\_REMODELLING\_OF\_PE, REACTOME\_ACYL\_CHAIN\_REMODELLING\_OF\_PE  
GOBP\_T\_HELPER\_CELL\_LINEAGE\_COMMITMENT, GOBP\_T\_HELPER\_CELL\_LINEAGE\_COMMITMENT  
WP\_HEART\_DEVELOPMENT, WP\_HEART\_DEVELOPMENT  
GOMF\_ION\_GATED\_CHANNEL\_ACTIVITY, GOMF\_ION\_GATED\_CHANNEL\_ACTIVITY  
HP\_FRONTOTEMPORAL\_CEREBRAL\_ATROPHY, HP\_FRONTOTEMPORAL\_CEREBRAL\_ATROPHY  
HP\_MALE\_INFERTILITY, HP\_MALE\_INFERTILITY  
HP\_RENAL\_CORTICOMEDULLARY\_CYSTS, HP\_RENAL\_CORTICOMEDULLARY\_CYSTS  
ODONNELL\_METASTASIS\_UP, ODONNELL\_METASTASIS\_UP  
HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_2ND\_FINGER, HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_2ND\_FINGER  
GOBP\_NUCLEAR\_MIGRATION, GOBP\_NUCLEAR\_MIGRATION  
GOBP\_NUCLEUS\_LOCALIZATION, GOBP\_NUCLEUS\_LOCALIZATION  
GOMF\_PHOSPHOLIPASE\_A1\_ACTIVITY, GOMF\_PHOSPHOLIPASE\_A1\_ACTIVITY  
GOBP\_NEGATIVE\_REGULATION\_OF\_OSTEOCLAST\_DIFFERENTIATION, GOBP\_NEGATIVE\_REGULATION\_OF\_OSTEOCLAST\_DIFFERENTIATION  
GOBP\_LENS\_FIBER\_CELL\_DEVELOPMENT, GOBP\_LENS\_FIBER\_CELL\_DEVELOPMENT  
GOBP\_POSITIVE\_REGULATION\_OF\_LIPOPROTEIN\_PARTICLE\_CLEARANCE, GOBP\_POSITIVE\_REGULATION\_OF\_LIPOPROTEIN\_PARTICLE\_CLEARANCE  
GOBP\_NEGATIVE\_REGULATION\_OF\_TISSUE\_REMODELING, GOBP\_NEGATIVE\_REGULATION\_OF\_TISSUE\_REMODELING  
GOMF\_1\_ACYL\_2\_LYSOPHOSPHATIDYLSERINE\_ACYLHYDROLASE\_ACTIVITY, GOMF\_1\_ACYL\_2\_LYSOPHOSPHATIDYLSERINE\_ACYLHYDROLASE\_ACTIVITY  
GOBP\_REGULATION\_OF\_VASCULAR\_ASSOCIATED\_SMOOTH\_MUSCLE\_CELL\_DEDIFFERENTIATION, GOBP\_REGULATION\_OF\_VASCULAR\_ASSOCIATED\_SMOOTH\_MUSCLE\_CELL\_DEDIFFERENTIATION  
GOBP\_MODIFICATION\_OF\_SYNAPTIC\_STRUCTURE, GOBP\_MODIFICATION\_OF\_SYNAPTIC\_STRUCTURE  
MIKKELSEN\_MEF\_LCP\_WITH\_H3K27ME3, MIKKELSEN\_MEF\_LCP\_WITH\_H3K27ME3  
GOCC\_STEREOCILIIUM\_TIP, GOCC\_STEREOCILIIUM\_TIP  
GOBP\_DIAPHRAGM\_DEVELOPMENT, GOBP\_DIAPHRAGM\_DEVELOPMENT  
GOCC\_NEUROFILAMENT, GOCC\_NEUROFILAMENT  
GOBP\_POSITIVE\_REGULATION\_OF\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION, GOBP\_POSITIVE\_REGULATION\_OF\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION  
GOBP\_POSITIVE\_REGULATION\_OF\_CARTILAGE\_DEVELOPMENT, GOBP\_POSITIVE\_REGULATION\_OF\_CARTILAGE\_DEVELOPMENT  
REACTOME\_PHOSPHOLIPASE\_C\_MEDIATED\_CASCADE\_FGFR2, REACTOME\_PHOSPHOLIPASE\_C\_MEDIATED\_CASCADE\_FGFR2  
GOBP\_CEREBELLAR\_GRANULAR\_LAYER\_FORMATION, GOBP\_CEREBELLAR\_GRANULAR\_LAYER\_FORMATION