

GSE42724\_MEMORY\_BCELL\_VS\_PLASMABLAST\_DN, GSE42724\_MEMORY\_BCELL\_VS\_PLASMABLAST\_DN  
GSE22919\_RESTING\_VS\_IL2\_IL12\_IL15\_STIM\_NK\_CELL\_DN, GSE22919\_RESTING\_VS\_IL2\_IL12\_IL15\_STIM\_NK\_CELL\_DN  
GSE46606\_IRF4\_KO\_VS\_WT\_UNSTIM\_BCELL\_UP, GSE46606\_IRF4\_KO\_VS\_WT\_UNSTIM\_BCELL\_UP  
GSE26343\_WT\_VS\_NFAT5\_KO\_MACROPHAGE\_UP, GSE26343\_WT\_VS\_NFAT5\_KO\_MACROPHAGE\_UP  
GSE24726\_WT\_VS\_E2\_2\_KO\_PDC\_DAY6\_POST\_DELETION\_DN, GSE24726\_WT\_VS\_E2\_2\_KO\_PDC\_DAY6\_POST\_DELETION\_DN  
GSE7348\_UNSTIM\_VS\_TOLERIZED\_AND\_LPS\_STIM\_MACROPHAGE\_UP, GSE7348\_UNSTIM\_VS\_TOLERIZED\_AND\_LPS\_STIM\_MACROPHAGE\_UP  
GSE23308\_WT\_VS\_MINERALCORTICOID\_REC\_KO\_MACROPHAGE\_UP, GSE23308\_WT\_VS\_MINERALCORTICOID\_REC\_KO\_MACROPHAGE\_UP  
GSE11864\_CSF1\_VS\_CSF1\_IFNG\_PAM3CYS\_IN\_MAC\_UP, GSE11864\_CSF1\_VS\_CSF1\_IFNG\_PAM3CYS\_IN\_MAC\_UP  
GSE17721\_LPS\_VS\_GARDIQUIMOD\_6H\_BMDC\_DN, GSE17721\_LPS\_VS\_GARDIQUIMOD\_6H\_BMDC\_DN  
GSE32986\_CURDLAN\_LOWDOSE\_VS\_CURDLAN\_HIGHDOSE\_STIM\_DC\_UP, GSE32986\_CURDLAN\_LOWDOSE\_VS\_CURDLAN\_HIGHDOSE\_STIM\_DC\_UP  
GSE42724\_NAIVE\_BCELL\_VS\_PLASMABLAST\_DN, GSE42724\_NAIVE\_BCELL\_VS\_PLASMABLAST\_DN  
GSE17721\_CTRL\_VS\_GARDIQUIMOD\_4H\_BMDC\_UP, GSE17721\_CTRL\_VS\_GARDIQUIMOD\_4H\_BMDC\_UP  
GSE10240\_IL17\_VS\_IL17\_AND\_IL22\_STIM\_PRIMARY\_BRONCHIAL\_EPITHELIAL\_CELLS\_UP, GSE10240\_IL17\_VS\_IL17\_AND\_IL22\_STIM\_PRIMARY\_BRONCHIAL\_EPITHELIAL\_CELLS\_UP  
GSE6092\_IFNG\_VS\_IFNG\_AND\_B\_BURGDORFERI\_INF\_ENDOTHELIAL\_CELL\_UP, GSE6092\_IFNG\_VS\_IFNG\_AND\_B\_BURGDORFERI\_INF\_ENDOTHELIAL\_CELL\_UP  
GSE24210\_CTRL\_VS\_IL35\_TREATED\_TCONV\_CD4\_TCELL\_UP, GSE24210\_CTRL\_VS\_IL35\_TREATED\_TCONV\_CD4\_TCELL\_UP  
GSE4590\_PRE\_BCELL\_VS\_LARGE\_PRE\_BCELL\_DN, GSE4590\_PRE\_BCELL\_VS\_LARGE\_PRE\_BCELL\_DN  
GSE9239\_CTRL\_VS\_TNF\_INHIBITOR\_TREATED\_DC\_UP, GSE9239\_CTRL\_VS\_TNF\_INHIBITOR\_TREATED\_DC\_UP  
GSE9037\_CTRL\_VS\_LPS\_4H\_STIM\_IRAK4\_KO\_BMDM\_UP, GSE9037\_CTRL\_VS\_LPS\_4H\_STIM\_IRAK4\_KO\_BMDM\_UP  
GSE22229\_RENAL\_TRANSPLANT\_VS\_HEALTHY\_PBMCDN, GSE22229\_RENAL\_TRANSPLANT\_VS\_HEALTHY\_PBMCDN  
GSE27786\_BCELL\_VS\_NKCELL\_DN, GSE27786\_BCELL\_VS\_NKCELL\_DN  
GSE13522\_CTRL\_VS\_T\_CRUZL\_Y\_STRAIN\_INF\_SKIN\_IFNAR\_KO\_UP, GSE13522\_CTRL\_VS\_T\_CRUZL\_Y\_STRAIN\_INF\_SKIN\_IFNAR\_KO\_UP  
MIR3190\_3P, MIR3190\_3P  
GSE17186\_MEMORY\_VS\_NAIVE\_BCELL\_UP, GSE17186\_MEMORY\_VS\_NAIVE\_BCELL\_UP  
MIR3920, MIR3920  
GSE39152\_CD103\_NEG\_VS\_POS\_MEMORY\_CD8\_TCELL\_UP, GSE39152\_CD103\_NEG\_VS\_POS\_MEMORY\_CD8\_TCELL\_UP  
BYSTRYKH\_HEMATOPOIESIS\_STEM\_CELL\_QTL\_CIS, BYSTRYKH\_HEMATOPOIESIS\_STEM\_CELL\_QTL\_CIS  
GSE8868\_SPLEEN\_VS\_INTESTINE\_CD11B\_POS\_CD11C\_NEG\_DC\_DN, GSE8868\_SPLEEN\_VS\_INTESTINE\_CD11B\_POS\_CD11C\_NEG\_DC\_DN  
GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_IL4\_STIM\_UP, GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_IL4\_STIM\_UP  
MIR2276\_3P, MIR2276\_3P  
MIR2115\_5P, MIR2115\_5P  
GOBP\_REGULATION\_OF\_PROTEIN\_BINDING, GOBP\_REGULATION\_OF\_PROTEIN\_BINDING  
HP\_CLEFT\_LIP, HP\_CLEFT\_LIP  
HP\_LOW\_ANTERIOR\_HAIRLINE, HP\_LOW\_ANTERIOR\_HAIRLINE  
MIR6509\_3P, MIR6509\_3P  
GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_ROSIGLITAZONE\_STIM\_UP, GSE25123\_WT\_VS\_PPARG\_KO\_MACROPHAGE\_ROSIGLITAZONE\_STIM\_UP  
GSE22282\_HYPOXIA\_VS\_NORMOXIA\_MYELOID\_DC\_DN, GSE22282\_HYPOXIA\_VS\_NORMOXIA\_MYELOID\_DC\_DN  
MIR10526\_3P, MIR10526\_3P  
MIR1285\_3P, MIR1285\_3P  
MIR3187\_5P, MIR3187\_5P  
GSE30083\_SP1\_VS\_SP3\_THYMOCYTE\_DN, GSE30083\_SP1\_VS\_SP3\_THYMOCYTE\_DN  
HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_EAR, HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_EAR  
GSE7852\_TREG\_VS\_TCONV\_LN\_DN, GSE7852\_TREG\_VS\_TCONV\_LN\_DN  
MIR6860, MIR6860  
PDGF\_ERK\_DN.V1\_UP, PDGF\_ERK\_DN.V1\_UP  
MIR5189\_5P, MIR5189\_5P  
MIR612, MIR612  
GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_HEXOSYL\_GROUPS, GOMF\_TRANSFERASE\_ACTIVITY\_TRANSFERRING\_HEXOSYL\_GROUPS  
GOBP\_PROTEIN\_O\_LINKED\_GLYCOSYLATION, GOBP\_PROTEIN\_O\_LINKED\_GLYCOSYLATION  
HP\_COARCTATION\_OF\_AORTA, HP\_COARCTATION\_OF\_AORTA  
GSE2770\_UNTREATED\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_2H\_UP, GSE2770\_UNTREATED\_VS\_TGFB\_AND\_IL12\_TREATED\_ACT\_CD4\_TCELL\_2H\_UP  
MIR1260A\_MIR1260B, MIR1260A\_MIR1260B  
MIR1915\_3P, MIR1915\_3P  
WP\_JOUBERT\_SYNDROME, WP\_JOUBERT\_SYNDROME  
MTOR\_UP.V1\_DN, MTOR\_UP.V1\_DN  
RUVBL2\_TARGET\_GENES, RUVBL2\_TARGET\_GENES  
HP\_PROMINENT\_NOSE, HP\_PROMINENT\_NOSE  
MIR2467\_5P, MIR2467\_5P  
GRE\_C, GRE\_C  
GOBP\_POSITIVE\_REGULATION\_OF\_AUTOPHAGY, GOBP\_POSITIVE\_REGULATION\_OF\_AUTOPHAGY  
GSE14415\_INDUCED\_TREG\_VS\_FOXP3\_KO\_INDUCED\_TREG\_IL2\_CULTURE\_DN, GSE14415\_INDUCED\_TREG\_VS\_FOXP3\_KO\_INDUCED\_TREG\_IL2\_CULTURE\_DN  
GOBP\_MANNOSYLATION, GOBP\_MANNOSYLATION  
GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_BINDING, GOBP\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_BINDING  
MIR4686, MIR4686  
ROSS\_ACUTE\_MYELOID\_LEUKEMIA\_CBF, ROSS\_ACUTE\_MYELOID\_LEUKEMIA\_CBF  
CHIANG\_LIVER\_CANCER\_SUBCLASS\_CTNNB1\_UP, CHIANG\_LIVER\_CANCER\_SUBCLASS\_CTNNB1\_UP  
MIR6758\_5P, MIR6758\_5P  
ZSCAN23\_TARGET\_GENES, ZSCAN23\_TARGET\_GENES  
MIR3120\_5P, MIR3120\_5P  
HP\_SHORT\_FOOT, HP\_SHORT\_FOOT  
GOMF\_PROTEIN\_SERINE\_PHOSPHATASE\_ACTIVITY, GOMF\_PROTEIN\_SERINE\_PHOSPHATASE\_ACTIVITY  
DHIMAN\_PBMCD\_ATTENUVAX\_AGE\_15\_25YO\_SUBQ\_7\_OR\_14DY\_UP, DHIMAN\_PBMCD\_ATTENUVAX\_AGE\_15\_25YO\_SUBQ\_7\_OR\_14DY\_UP  
MIR550B\_3P, MIR550B\_3P  
ATGTCAC\_MIR489, ATGTCAC\_MIR489  
DAVICIONI\_PAX\_FOXO1\_SIGNATURE\_IN\_ARMS\_UP, DAVICIONI\_PAX\_FOXO1\_SIGNATURE\_IN\_ARMS\_UP  
GTAAGAT\_MIR200A, GTAAGAT\_MIR200A  
GOBP\_POSITIVE\_REGULATION\_OF\_MITOCHONDRIAL\_FISSION, GOBP\_POSITIVE\_REGULATION\_OF\_MITOCHONDRIAL\_FISSION  
HP\_ABNORMAL\_ERYTHROCYTE\_ENZYME\_LEVEL, HP\_ABNORMAL\_ERYTHROCYTE\_ENZYME\_LEVEL  
GSE21063\_CTRL\_VS\_ANTLJGM\_STIM\_BCELL\_8H\_DN, GSE21063\_CTRL\_VS\_ANTLJGM\_STIM\_BCELL\_8H\_DN  
GOBP\_NEGATIVE\_REGULATION\_OF\_DNA\_REPAIR, GOBP\_NEGATIVE\_REGULATION\_OF\_DNA\_REPAIR  
GOBP\_REGULATION\_OF\_CHROMATIN\_BINDING, GOBP\_REGULATION\_OF\_CHROMATIN\_BINDING  
MIR5704, MIR5704  
GOMF\_MANNOSYLTRANSFERASE\_ACTIVITY, GOMF\_MANNOSYLTRANSFERASE\_ACTIVITY  
MIR1272, MIR1272  
HP\_ABNORMALITY\_OF\_MIDBRAIN\_MORPHOLOGY, HP\_ABNORMALITY\_OF\_MIDBRAIN\_MORPHOLOGY  
GOBP\_PROTEOGLYCAN\_METABOLIC\_PROCESS, GOBP\_PROTEOGLYCAN\_METABOLIC\_PROCESS  
KEGG\_GLYCOSAMINOGLYCAN\_BIOSYNTHESIS\_CHONDROITIN\_SULFATE, KEGG\_GLYCOSAMINOGLYCAN\_BIOSYNTHESIS\_CHONDROITIN\_SULFATE  
MIR15A\_3P, MIR15A\_3P  
MIR379\_5P, MIR379\_5P  
NIKOLSKY\_BREAST\_CANCER\_6P24\_P22\_AMPLICON, NIKOLSKY\_BREAST\_CANCER\_6P24\_P22\_AMPLICON  
HP\_ABNORMALITY\_OF\_THE\_COLUMELLA, HP\_ABNORMALITY\_OF\_THE\_COLUMELLA  
HP\_BIPARIETAL\_NARROWING, HP\_BIPARIETAL\_NARROWING  
GOBP\_PEPTIDYL\_TYROSINE\_AUTOPHOSPHORYLATION, GOBP\_PEPTIDYL\_TYROSINE\_AUTOPHOSPHORYLATION  
GOBP\_PROTEOGLYCAN\_BIOSYNTHETIC\_PROCESS, GOBP\_PROTEOGLYCAN\_BIOSYNTHETIC\_PROCESS  
HP\_DEVELOPMENTAL\_GLAUCOMA, HP\_DEVELOPMENTAL\_GLAUCOMA  
GSE40274\_CTRL\_VS\_FOXP3\_AND\_SATB1\_TRANSDUCED\_ACTIVATED\_CD4\_TCELL\_DN, GSE40274\_CTRL\_VS\_FOXP3\_AND\_SATB1\_TRANSDUCED\_ACTIVATED\_CD4\_TCELL\_DN  
GOBP\_FUCOSYLATION, GOBP\_FUCOSYLATION  
HP\_IMPAIRED\_VISUOSPATIAL\_CONSTRUCTIVE\_COGNITION, HP\_IMPAIRED\_VISUOSPATIAL\_CONSTRUCTIVE\_COGNITION  
KEGG\_GLYCOPHINGOLIPID\_BIOSYNTHESIS\_GLOBO\_SERIES, KEGG\_GLYCOPHINGOLIPID\_BIOSYNTHESIS\_GLOBO\_SERIES  
PPARG\_01, PPARG\_01  
HP\_ABNORMALITY\_OF\_THE\_PHALANGES\_OF\_THE\_HALLUX, HP\_ABNORMALITY\_OF\_THE\_PHALANGES\_OF\_THE\_HALLUX  
GOBP\_INOSITOL\_PHOSPHATE\_METABOLIC\_PROCESS, GOBP\_INOSITOL\_PHOSPHATE\_METABOLIC\_PROCESS  
GOBP\_ENDOCARDIAL\_CUSHION\_DEVELOPMENT, GOBP\_ENDOCARDIAL\_CUSHION\_DEVELOPMENT  
GOMF\_LYSOPHOSPHOLIPID\_ACYLTRANSFERASE\_ACTIVITY, GOMF\_LYSOPHOSPHOLIPID\_ACYLTRANSFERASE\_ACTIVITY  
GOBP\_CLEAVAGE\_FURROW\_FORMATION, GOBP\_CLEAVAGE\_FURROW\_FORMATION  
REACTOME\_DOWNREGULATION\_OF\_ERBB2\_SIGNALING, REACTOME\_DOWNREGULATION\_OF\_ERBB2\_SIGNALING  
HP\_RETINAL\_COLOBOMA, HP\_RETINAL\_COLOBOMA  
WP\_WNT\_SIGNALING\_PATHWAY\_NETPATH, WP\_WNT\_SIGNALING\_PATHWAY\_NETPATH  
GOBP\_RESPONSE\_TO\_ISCHEMIA, GOBP\_RESPONSE\_TO\_ISCHEMIA  
GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_KINASE\_B\_SIGNALING, GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_KINASE\_B\_SIGNALING  
GOBP\_POLYOL\_CATABOLIC\_PROCESS, GOBP\_POLYOL\_CATABOLIC\_PROCESS  
REACTOME\_TRANSPORT\_OF\_NUCLEOTIDE\_SUGARS, REACTOME\_TRANSPORT\_OF\_NUCLEOTIDE\_SUGARS  
GAVIN\_IL2\_RESPONSIVE\_FOXP3\_TARGETS\_UP, GAVIN\_IL2\_RESPONSIVE\_FOXP3\_TARGETS\_UP  
REACTOME\_ELEVATION\_OF\_CYTOSOLIC\_CA2\_LEVELS, REACTOME\_ELEVATION\_OF\_CYTOSOLIC\_CA2\_LEVELS  
GSE13762\_CTRL\_VS\_125\_VITAMIND\_DAYS\_DC\_DN, GSE13762\_CTRL\_VS\_125\_VITAMIND\_DAYS\_DC\_DN  
GSE40274\_FOXP3\_VS\_FOXP3\_AND\_HELIOS\_TRANSDUCED\_ACTIVATED\_CD4\_TCELL\_UP, GSE40274\_FOXP3\_VS\_FOXP3\_AND\_HELIOS\_TRANSDUCED\_ACTIVATED\_CD4\_TCELL\_UP  
GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_SUBSTRATE\_JUNCTION\_ORGANIZATION, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_SUBSTRATE\_JUNCTION\_ORGANIZATION  
GOBP\_HISTONE\_H3\_K36\_DEMETHYLATION, GOBP\_HISTONE\_H3\_K36\_DEMETHYLATION  
GOMF\_RACEMASE\_AND\_EPIMERASE\_ACTIVITY\_ACTING\_ON\_CARBOHYDRATES\_AND\_DERIVATIVES, GOMF\_RACEMASE\_AND\_EPIMERASE\_ACTIVITY\_ACTING\_ON\_CARBOHYDRATES\_AND\_DERIVATIVES  
GOMF\_HISTONE\_DEMETHYLASE\_ACTIVITY\_H3\_K36\_SPECIFIC, GOMF\_HISTONE\_DEMETHYLASE\_ACTIVITY\_H3\_K36\_SPECIFIC  
GOBP\_DERMATAN\_SULFATE\_PROTEOGLYCAN\_METABOLIC\_PROCESS, GOBP\_DERMATAN\_SULFATE\_PROTEOGLYCAN\_METABOLIC\_PROCESS  
GOBP\_INSULIN\_METABOLIC\_PROCESS, GOBP\_INSULIN\_METABOLIC\_PROCESS  
GOBP\_LOCOMOTION\_INVOLVED\_IN\_LOCOMOTORY\_BEHAVIOR, GOBP\_LOCOMOTION\_INVOLVED\_IN\_LOCOMOTORY\_BEHAVIOR  
HP\_SHORT\_COLUMELLA, HP\_SHORT\_COLUMELLA  
MIR8056, MIR8056  
WP\_HEART\_DEVELOPMENT, WP\_HEART\_DEVELOPMENT  
GSE29618\_PRE\_VS\_DAY7\_FLU\_VACCINE\_PDC\_UP, GSE29618\_PRE\_VS\_DAY7\_FLU\_VACCINE\_PDC\_UP  
GSE45365\_NK\_CELL\_VS\_CD11B\_DC\_UP, GSE45365\_NK\_CELL\_VS\_CD11B\_DC\_UP  
GOBP\_STORE\_OPERATED\_CALCIIUM\_ENTRY, GOBP\_STORE\_OPERATED\_CALCIIUM\_ENTRY  
GOBP\_ADHERENS\_JUNCTION\_MAINTENANCE, GOBP\_ADHERENS\_JUNCTION\_MAINTENANCE  
HP\_ABNORMALITY\_OF\_THE\_CRUS\_OF\_THE\_HELIX, HP\_ABNORMALITY\_OF\_THE\_CRUS\_OF\_THE\_HELIX  
HP\_SLOW\_PUPILLARY\_LIGHT\_RESPONSE, HP\_SLOW\_PUPILLARY\_LIGHT\_RESPONSE  
REACTOME\_REPRESSION\_OF\_WNT\_TARGET\_GENES, REACTOME\_REPRESSION\_OF\_WNT\_TARGET\_GENES  
HP\_ABNORMAL\_RIGHT\_VENTRICULAR\_FUNCTION, HP\_ABNORMAL\_RIGHT\_VENTRICULAR\_FUNCTION  
GOMF\_INOSITOL\_PHOSPHATE\_PHOSPHATASE\_ACTIVITY, GOMF\_INOSITOL\_PHOSPHATE\_PHOSPHATASE\_ACTIVITY  
REACTOME\_SYNTHESIS\_OF\_PIP3\_AT\_THE\_GOLGI\_MEMBRANE, REACTOME\_SYNTHESIS\_OF\_PIP3\_AT\_THE\_GOLGI\_MEMBRANE  
GOBP\_SPERM\_AXONEME\_ASSEMBLY, GOBP\_SPERM\_AXONEME\_ASSEMBLY  
HP\_RECURRENT\_INFECTION\_OF\_THE\_GASTROINTESTINAL\_TRACT, HP\_RECURRENT\_INFECTION\_OF\_THE\_GASTROINTESTINAL\_TRACT  
HP\_WIDELY\_SPACED\_MAXILLARY\_CENTRAL\_INCISORS, HP\_WIDELY\_SPACED\_MAXILLARY\_CENTRAL\_INCISORS  
GSE16450\_CTRL\_VS\_IFNA\_12H\_STIM\_MATURE\_NEURON\_CELL\_LINE\_UP, GSE16450\_CTRL\_VS\_IFNA\_12H\_STIM\_MATURE\_NEURON\_CELL\_LINE\_UP  
GCAAGAC\_MIR431, GCAAGAC\_MIR431  
GOBP\_REGULATION\_OF\_NON\_MEMBRANE\_SPANNING\_PROTEIN\_TYROSINE\_KINASE\_ACTIVITY, GOBP\_REGULATION\_OF\_NON\_MEMBRANE\_SPANNING\_PROTEIN\_TYROSINE\_KINASE\_ACTIVITY  
HOFT\_CD4\_POSITIVE\_ALPHA\_BETA\_MEMORY\_T\_CELL\_BCG\_VACCINE\_AGE\_18\_45YO\_7DY\_UP, HOFT\_CD4\_POSITIVE\_ALPHA\_BETA\_MEMORY\_T\_CELL\_BCG\_VACCINE\_AGE\_18\_45YO\_7DY\_UP  
GOBP\_BLOOD\_VESSEL\_REMODELING, GOBP\_BLOOD\_VESSEL\_REMODELING  
REACTOME\_SIGNALING\_BY\_BMP, REACTOME\_SIGNALING\_BY\_BMP  
GOBP\_PHAGOLYSOSOME\_ASSEMBLY, GOBP\_PHAGOLYSOSOME\_ASSEMBLY  
GOMF\_BHLH\_TRANSCRIPTION\_FACTOR\_BINDING, GOMF\_BHLH\_TRANSCRIPTION\_FACTOR\_BINDING  
GOBP\_RETINA\_VASCULATURE\_DEVELOPMENT\_IN\_CAMERA\_TYPE\_EYE, GOBP\_RETINA\_VASCULATURE\_DEVELOPMENT\_IN\_CAMERA\_TYPE\_EYE  
GOMF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVITY, GOMF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_SERINE\_THREONINE\_KINASE\_ACTIVITY

WT\_CD40L\_IL2\_IL5\_DAY1\_STIMULATED\_BCELL\_UP, GSE46606\_IRF4HIGH\_VS\_WT\_CD40L\_IL2\_IL5\_DAY1\_STIMULATED\_BCELL\_UP