

GSE4142\_NAIVE\_BCELL\_UP,WP\_PLASMA\_CELL\_UP,GSE4142\_NAIVE\_BCELL\_UP,WP\_PLASMA\_CELL\_UP  
GSE8921\_UNSTIM\_VS\_TLRI\_2\_STIM\_MONOCYTE\_3H\_DN,GSE8921\_UNSTIM\_VS\_TLRI\_2\_STIM\_MONOCYTE\_3H\_DN  
GSE39820\_TGFBETA1\_VS\_TGFBETA3\_IN\_IL6\_TREATED\_CD4\_TCELL\_UP,GSE39820\_TGFBETA1\_VS\_TGFBETA3\_IN\_IL6\_TREATED\_CD4\_TCELL\_UP  
GSE17186\_MEMORY\_VS\_CD21HIGH\_TRANSITIONAL\_BCELL\_DN,GSE17186\_MEMORY\_VS\_CD21HIGH\_TRANSITIONAL\_BCELL\_DN  
GSE16522\_MEMORY\_VS\_NAIVE\_ANTLCD3CD28\_STIM\_CD8\_TCELL\_UP,GSE16522\_MEMORY\_VS\_NAIVE\_ANTLCD3CD28\_STIM\_CD8\_TCELL\_UP  
REACTOME\_RAC1\_GTPASE\_CYCLE,REACTOME\_RAC1\_GTPASE\_CYCLE  
GSE39820\_CTRL\_VS\_TGFBETA1\_IL6\_CD4\_TCELL\_DN,GSE39820\_CTRL\_VS\_TGFBETA1\_IL6\_CD4\_TCELL\_DN  
GSE39820\_CTRL\_VS\_TGFBETA3\_IL6\_CD4\_TCELL\_DN,GSE39820\_CTRL\_VS\_TGFBETA3\_IL6\_CD4\_TCELL\_DN  
GSE36392\_TYPE\_2\_MYELOID\_VS\_NEUTROPHIL\_IL25\_TREATED\_LUNG\_UP,GSE36392\_TYPE\_2\_MYELOID\_VS\_NEUTROPHIL\_IL25\_TREATED\_LUNG\_UP  
GSE15330\_HSC\_VS\_MEGAKARYOCYTE\_ERYTHROID\_PROGENITOR\_IKAROS\_KO\_UP,GSE15330\_HSC\_VS\_MEGAKARYOCYTE\_ERYTHROID\_PROGENITOR\_IKAROS\_KO\_UP  
GSE41867\_NAIVE\_VS\_DAY8\_LCMV\_CLONE13\_EFFECTOR\_CD8\_TCELL\_UP,GSE41867\_NAIVE\_VS\_DAY8\_LCMV\_CLONE13\_EFFECTOR\_CD8\_TCELL\_UP  
GSE17721\_0\_5H\_VS\_4H\_POLYIC\_BMDC\_DN,GSE17721\_0\_5H\_VS\_4H\_POLYIC\_BMDC\_DN  
GSE16451\_IMMATURE\_VS\_MATURE\_NEURON\_CELL\_LINE\_WEST\_EQUINE\_ENC\_VIRUS\_UP,GSE16451\_IMMATURE\_VS\_MATURE\_NEURON\_CELL\_LINE\_WEST\_EQUINE\_ENC\_VIRUS\_UP  
GSE32255\_UNSTIM\_VS\_4H\_LPS\_STIM\_DC\_UP,GSE32255\_UNSTIM\_VS\_4H\_LPS\_STIM\_DC\_UP  
GSE15330\_LYMPHOID\_MULTIPOTENT\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_IKAROS\_KO\_DN,GSE15330\_LYMPHOID\_MULTIPOTENT\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_IKAROS\_KO\_DN  
GSE17721\_POLYIC\_VS\_CPG\_0\_5H\_BMDC\_DN,GSE17721\_POLYIC\_VS\_CPG\_0\_5H\_BMDC\_DN  
REACTOME\_CDC42\_GTPASE\_CYCLE,REACTOME\_CDC42\_GTPASE\_CYCLE  
WP\_INTEGRATED\_BREAST\_CANCER\_PATHWAY,WP\_INTEGRATED\_BREAST\_CANCER\_PATHWAY  
MIR6807\_3P,MIR6807\_3P  
GSE19941\_IL10\_KO\_VS\_IL10\_KO\_AND\_NFKBP50\_KO\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_UP,GSE19941\_IL10\_KO\_VS\_IL10\_KO\_AND\_NFKBP50\_KO\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_UP  
GSE37301\_HEMATOPOIETIC\_STEM\_CELL\_VS\_LYMPHOID\_PRIMED\_MPP\_UP,GSE37301\_HEMATOPOIETIC\_STEM\_CELL\_VS\_LYMPHOID\_PRIMED\_MPP\_UP  
IKEDA\_MIR30\_TARGETS\_UP,IKEDA\_MIR30\_TARGETS\_UP  
MIR421,MIR421  
GOBP\_TRANSPORT\_ALONG\_MICROTUBULE,GOBP\_TRANSPORT\_ALONG\_MICROTUBULE  
GOBP\_CYTOSKELETON\_DEPENDENT\_INTRACELLULAR\_TRANSPORT,GOBP\_CYTOSKELETON\_DEPENDENT\_INTRACELLULAR\_TRANSPORT  
ACACTAC\_MIR1423P,ACACTAC\_MIR1423P  
GSE15330\_HSC\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_IKAROS\_KO\_UP,GSE15330\_HSC\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_IKAROS\_KO\_UP  
GSE19941\_LPS\_VS\_LPS\_AND\_IL10\_STIM\_IL10\_KO\_NFKBP50\_KO\_MACROPHAGE\_DN,GSE19941\_LPS\_VS\_LPS\_AND\_IL10\_STIM\_IL10\_KO\_NFKBP50\_KO\_MACROPHAGE\_DN  
GOBP\_MICROTUBULE\_BASED\_TRANSPORT,GOBP\_MICROTUBULE\_BASED\_TRANSPORT  
CTACTGT\_MIR199A,CTACTGT\_MIR199A  
MIR299\_5P,MIR299\_5P  
MIR2053,MIR2053  
GSE17721\_12H\_VS\_24H\_GARDIQUIMOD\_BMDC\_UP,GSE17721\_12H\_VS\_24H\_GARDIQUIMOD\_BMDC\_UP  
GSE19941\_IL10\_KO\_VS\_IL10\_KO\_AND\_NFKBP50\_KO\_UNSTIM\_MACROPHAGE\_DN,GSE19941\_IL10\_KO\_VS\_IL10\_KO\_AND\_NFKBP50\_KO\_UNSTIM\_MACROPHAGE\_DN  
GSE22025\_PROGESTERONE\_VS\_TGFB1\_AND\_PROGESTERONE\_TREATED\_CD4\_TCELL\_DN,GSE22025\_PROGESTERONE\_VS\_TGFB1\_AND\_PROGESTERONE\_TREATED\_CD4\_TCELL\_DN  
GSE25087\_FETAL\_VS\_ADULT\_TREG\_DN,GSE25087\_FETAL\_VS\_ADULT\_TREG\_DN  
MIR8087,MIR8087  
GSE37605\_NOD\_VS\_C57BL6\_IRES\_GFP\_TREG\_UP,GSE37605\_NOD\_VS\_C57BL6\_IRES\_GFP\_TREG\_UP  
MIR891A\_3P,MIR891A\_3P  
GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_4H\_BMDC\_DN,GSE17721\_PAM3CSK4\_VS\_GADIQUIMOD\_4H\_BMDC\_DN  
MIR2054,MIR2054  
CTCTGGA\_MIR520A\_MIR525,CTCTGGA\_MIR520A\_MIR525  
WP\_PROLACTIN\_SIGNALING\_PATHWAY,WP\_PROLACTIN\_SIGNALING\_PATHWAY  
KEGG\_FC\_GAMMA\_R\_MEDIATED\_PHAGOCYTOSIS,KEGG\_FC\_GAMMA\_R\_MEDIATED\_PHAGOCYTOSIS  
KEGG\_FOCAL\_ADHESION,KEGG\_FOCAL\_ADHESION  
HP\_HYPERLORDOSIS,HP\_HYPERLORDOSIS  
ATGCAGT\_MIR217,ATGCAGT\_MIR217  
GSE6875\_TCONV\_VS\_TREG\_UP,GSE6875\_TCONV\_VS\_TREG\_UP  
WP\_REGULATION\_OF\_ACTIN\_CYTOSKELETON,WP\_REGULATION\_OF\_ACTIN\_CYTOSKELETON  
MIR3190\_5P,MIR3190\_5P  
HP\_CRANIOSYNOSTOSIS,HP\_CRANIOSYNOSTOSIS  
COULOUARN\_TEMPORAL\_TGFB1\_SIGNATURE\_UP,COULOUARN\_TEMPORAL\_TGFB1\_SIGNATURE\_UP  
MIR651\_5P,MIR651\_5P  
WP\_BREAST\_CANCER\_PATHWAY,WP\_BREAST\_CANCER\_PATHWAY  
GOBP\_PEPTIDYL\_THREONINE\_MODIFICATION,GOBP\_PEPTIDYL\_THREONINE\_MODIFICATION  
MIR4423\_3P,MIR4423\_3P  
MIR6792\_5P,MIR6792\_5P  
WP\_RAS\_SIGNALING,WP\_RAS\_SIGNALING  
GGGNNTTCC\_NFKB\_Q6\_01,GGGNNTTCC\_NFKB\_Q6\_01  
MIR3168,MIR3168  
GOBP\_NEURON\_PROJECTION\_EXTENSION,GOBP\_NEURON\_PROJECTION\_EXTENSION  
KIM\_MYC\_AMPLIFICATION\_TARGETS\_DN,KIM\_MYC\_AMPLIFICATION\_TARGETS\_DN  
JECHLINGER\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_DN,JECHLINGER\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_DN  
CSE33513\_TCF7\_KO\_VS\_HET\_EARLY\_THYMIC\_PROGENITOR\_UP,CSE33513\_TCF7\_KO\_VS\_HET\_EARLY\_THYMIC\_PROGENITOR\_UP  
MIR942\_3P,MIR942\_3P  
GOBP\_ENDOPLASMIC\_RETICULUM\_ORGANIZATION,GOBP\_ENDOPLASMIC\_RETICULUM\_ORGANIZATION  
HP\_ABNORMAL\_AORTIC\_VALVE\_PHYSIOLOGY,HP\_ABNORMAL\_AORTIC\_VALVE\_PHYSIOLOGY  
HP\_CLONUS,HP\_CLONUS  
GOBP\_CELL\_SUBSTRATE\_JUNCTION\_ORGANIZATION,GOBP\_CELL\_SUBSTRATE\_JUNCTION\_ORGANIZATION  
GSE12198\_CTRL\_VS\_HIGH\_IL2\_STIM\_NK\_CELL\_UP,GSE12198\_CTRL\_VS\_HIGH\_IL2\_STIM\_NK\_CELL\_UP  
CHEBOTAEV\_GR\_TARGETS\_UP,CHEBOTAEV\_GR\_TARGETS\_UP  
GSE28783\_ANTL\_MIR33\_VS\_CTRL\_ATHEROSCLEROSIS\_MACROPHAGE\_DN,GSE28783\_ANTL\_MIR33\_VS\_CTRL\_ATHEROSCLEROSIS\_MACROPHAGE\_DN  
GSE16450\_CTRL\_VS\_IFNA\_6H\_STIM\_MATURE\_NEURON\_CELL\_LINE\_UP,GSE16450\_CTRL\_VS\_IFNA\_6H\_STIM\_MATURE\_NEURON\_CELL\_LINE\_UP  
MIR297,MIR297  
GOMF\_PROTEIN\_TYROSINE\_KINASE\_BINDING,GOMF\_PROTEIN\_TYROSINE\_KINASE\_BINDING  
GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_POLYMERIZATION,GOBP\_POSITIVE\_REGULATION\_OF\_PROTEIN\_POLYMERIZATION  
MIR5187\_3P,MIR5187\_3P  
GSE4142\_NAIVE\_VS\_MEMORY\_BCELL\_UP,GSE4142\_NAIVE\_VS\_MEMORY\_BCELL\_UP  
GOMF\_RECEPTOR\_TYROSINE\_KINASE\_BINDING,GOMF\_RECEPTOR\_TYROSINE\_KINASE\_BINDING  
GOMF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_KINASE\_ACTIVITY,GOMF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_KINASE\_ACTIVITY  
HFB3\_01,HFB3\_01  
GSE22229\_UNTREATED\_VS\_IMMUNOSUPP\_THERAPY\_RENAL\_TRANSPLANT\_PATIENT\_PBMC\_UP,GSE22229\_UNTREATED\_VS\_IMMUNOSUPP\_THERAPY\_RENAL\_TRANSPLANT\_PATIENT\_PBMC\_UP  
GOBP\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_POLYMERIZATION,GOBP\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_POLYMERIZATION  
PID\_FGF\_PATHWAY,PID\_FGF\_PATHWAY  
HP\_ABNORMAL\_LYMPHOCYTE\_MORPHOLOGY,HP\_ABNORMAL\_LYMPHOCYTE\_MORPHOLOGY  
MIR597\_5P,MIR597\_5P  
GSE22611\_NOD2\_TRANSD\_VS\_CTRL\_TRANSD\_HEK293\_MDP\_STIM\_2H\_UP,GSE22611\_NOD2\_TRANSD\_VS\_CTRL\_TRANSD\_HEK293\_MDP\_STIM\_2H\_UP  
MIR4670\_3P,MIR4670\_3P  
REACTOME\_SIGNALING\_BY\_FGFR,REACTOME\_SIGNALING\_BY\_FGFR  
HP\_LEUKOCYTOSIS,HP\_LEUKOCYTOSIS  
WP\_OSTEOBLAST\_DIFFERENTIATION,WP\_OSTEOBLAST\_DIFFERENTIATION  
GSE3994\_WT\_VS\_PAC1\_KO\_ACTIVATED\_MAST\_CELL\_UP,GSE3994\_WT\_VS\_PAC1\_KO\_ACTIVATED\_MAST\_CELL\_UP  
GSE21670\_UNTREATED\_VS\_TGFR\_TREATED\_STAT3\_KO\_CD4\_TCELL\_UP,GSE21670\_UNTREATED\_VS\_TGFR\_TREATED\_STAT3\_KO\_CD4\_TCELL\_UP  
GSE22611\_MUTANT\_NOD2\_TRANSDUCED\_VS\_CTRL\_HEK293T\_STIMULATED\_WITH\_MDP\_6H\_UP,GSE22611\_MUTANT\_NOD2\_TRANSDUCED\_VS\_CTRL\_HEK293T\_STIMULATED\_WITH\_MDP\_6H\_UP  
GOBP\_CARDIAC\_SEPTUM\_MORPHOGENESIS,GOBP\_CARDIAC\_SEPTUM\_MORPHOGENESIS  
REACTOME\_RND2\_GTPASE\_CYCLE,REACTOME\_RND2\_GTPASE\_CYCLE  
GOBP\_BONE\_MINERALIZATION,GOBP\_BONE\_MINERALIZATION  
GSE17721\_POLYIC\_VS\_PAM3CSK4\_2H\_BMDC\_DN,GSE17721\_POLYIC\_VS\_PAM3CSK4\_2H\_BMDC\_DN  
GOBP\_VENTRICULAR\_SEPTUM\_DEVELOPMENT,GOBP\_VENTRICULAR\_SEPTUM\_DEVELOPMENT  
MIR6081,MIR6081  
GOBP\_REGULATION\_OF\_EXTENT\_OF\_CELL\_GROWTH,GOBP\_REGULATION\_OF\_EXTENT\_OF\_CELL\_GROWTH  
GOBP\_CELLULAR\_RESPONSE\_TO\_ALCOHOL,GOBP\_CELLULAR\_RESPONSE\_TO\_ALCOHOL  
GSE19941\_UNSTIM\_VS\_LPS\_STIM\_IL10\_KO\_MACROPHAGE\_UP,GSE19941\_UNSTIM\_VS\_LPS\_STIM\_IL10\_KO\_MACROPHAGE\_UP  
WP\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_IN\_COLORECTAL\_CANCER,WP\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_IN\_COLORECTAL\_CANCER  
GOBP\_BIOMINERALIZATION,GOBP\_BIOMINERALIZATION  
GOBP\_MORPHOGENESIS\_OF\_A\_BRANCHING\_STRUCTURE,GOBP\_MORPHOGENESIS\_OF\_A\_BRANCHING\_STRUCTURE  
NAKAYA\_PBMCFUAD\_MALE\_AGE\_14\_27YO\_ID\_POSTBOOST\_VS\_ID\_PREIMM\_MF59\_ADJUVANTED\_IDY\_GENES\_IN\_BTMM40\_AND\_M53\_DN,NAKAYA\_PBMCFUAD\_MALE\_AGE\_14\_27YO\_ID\_POSTBOOST\_VS\_ID\_PREIMM\_MF59\_ADJUVANTED\_IDY\_GENES\_IN\_BTMM40\_AND\_M53\_DN  
HP\_ABNORMAL\_LEVELS\_OF\_ALPHA\_FETOPROTEIN,HP\_ABNORMAL\_LEVELS\_OF\_ALPHA\_FETOPROTEIN  
PID\_AVB3\_OPN\_PATHWAY,PID\_AVB3\_OPN\_PATHWAY  
PID\_EPHB\_FWD\_PATHWAY,PID\_EPHB\_FWD\_PATHWAY  
GOBP\_HISTONE\_H3\_K9\_MODIFICATION,GOBP\_HISTONE\_H3\_K9\_MODIFICATION  
GOBP\_EMBRYONIC\_APPENDAGE\_MORPHOGENESIS,GOBP\_EMBRYONIC\_APPENDAGE\_MORPHOGENESIS  
GOBP\_REGULATION\_OF\_BIOMINERALIZATION,GOBP\_REGULATION\_OF\_BIOMINERALIZATION  
GOBP\_PULMONARY\_VALVE\_DEVELOPMENT,GOBP\_PULMONARY\_VALVE\_DEVELOPMENT  
HP\_MICRODONTIA,HP\_MICRODONTIA  
GOBP\_APPENDAGE\_MORPHOGENESIS,GOBP\_APPENDAGE\_MORPHOGENESIS  
GSE43955\_1H\_VS\_42H\_ACT\_CD4\_TCELL\_DN,GSE43955\_1H\_VS\_42H\_ACT\_CD4\_TCELL\_DN  
HP\_DEFORMED\_SELLA\_TURCICA,HP\_DEFORMED\_SELLA\_TURCICA  
GOBP\_APPENDAGE\_DEVELOPMENT,GOBP\_APPENDAGE\_DEVELOPMENT  
REACTOME\_FGFR1\_MUTANT\_RECEPTOR\_ACTIVATION,REACTOME\_FGFR1\_MUTANT\_RECEPTOR\_ACTIVATION  
GSE5589\_WT\_VS\_IL10\_KO\_LPS\_STIM\_MACROPHAGE\_180MIN\_DN,GSE5589\_WT\_VS\_IL10\_KO\_LPS\_STIM\_MACROPHAGE\_180MIN\_DN  
WP\_HAIR\_FOLLICLE\_DEVELOPMENT\_ORGANOGENESIS\_PART\_2\_OF\_3,WP\_HAIR\_FOLLICLE\_DEVELOPMENT\_ORGANOGENESIS\_PART\_2\_OF\_3  
GOBP\_SPECIFICATION\_OF\_SYMMETRY,GOBP\_SPECIFICATION\_OF\_SYMMETRY  
GSE22935\_UNSTIM\_VS\_48H\_MBOVIS\_BCG\_STIM\_MYD88\_KO\_MACROPHAGE\_DN,GSE22935\_UNSTIM\_VS\_48H\_MBOVIS\_BCG\_STIM\_MYD88\_KO\_MACROPHAGE\_DN  
TOMLINS\_PROSTATE\_CANCER\_DN,TOMLINS\_PROSTATE\_CANCER\_DN  
PID\_NCADHERIN\_PATHWAY,PID\_NCADHERIN\_PATHWAY  
REACTOME\_DOWNSTREAM\_SIGNALING\_OF\_ACTIVATED\_FGFR1,REACTOME\_DOWNSTREAM\_SIGNALING\_OF\_ACTIVATED\_FGFR1  
GOBP\_ATRIOVENTRICULAR\_VALVE\_DEVELOPMENT,GOBP\_ATRIOVENTRICULAR\_VALVE\_DEVELOPMENT  
GOBP\_ENDOCARDIAL\_CUSHION\_DEVELOPMENT,GOBP\_ENDOCARDIAL\_CUSHION\_DEVELOPMENT  
MARTIN\_VIRAL\_GPCR\_SIGNALING\_DN,MARTIN\_VIRAL\_GPCR\_SIGNALING\_DN  
GOMF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_TYROSINE\_KINASE\_ADAPTOR\_ACTIVITY,GOMF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_TYROSINE\_KINASE\_ADAPTOR\_ACTIVITY  
ICHIBA\_GRAFT\_VERSUS\_HOST\_DISEASE\_3SD\_DN,ICHIBA\_GRAFT\_VERSUS\_HOST\_DISEASE\_3SD\_DN  
TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_DUCTAL\_NORMAL\_UP,TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_DUCTAL\_NORMAL\_UP  
HP\_PROMINENT\_NASAL\_TIP,HP\_PROMINENT\_NASAL\_TIP  
REACTOME\_IRS\_MEDIATED\_SIGNALLING,REACTOME\_IRS\_MEDIATED\_SIGNALLING  
GOBP\_CARDIOCYTE\_DIFFERENTIATION,GOBP\_CARDIOCYTE\_DIFFERENTIATION  
REACTOME\_INSULIN\_RECEPTOR\_SIGNALLING\_CASCADE,REACTOME\_INSULIN\_RECEPTOR\_SIGNALLING\_CASCADE  
MODULE\_202,MODULE\_202  
GOBP\_OUTFLOW\_TRACT\_MORPHOGENESIS,GOBP\_OUTFLOW\_TRACT\_MORPHOGENESIS  
GOBP\_POSITIVE\_REGULATION\_OF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_SERINE\_THREONINE\_KINASE\_SIGNALING\_PATHWAY,GOBP\_POSITIVE\_REGULATION\_OF\_TRANSMEMBRANE\_RECEPTOR\_PROTEIN\_SERINE\_THREONINE\_KINASE\_SIGNALING\_PATHWAY  
INGRAM\_SHH\_TARGETS\_DN,INGRAM\_SHH\_TARGETS\_DN  
GOBP\_ENDOCRINE\_SYSTEM\_DEVELOPMENT,GOBP\_ENDOCRINE\_SYSTEM\_DEVELOPMENT  
PID\_EPHA2\_FWD\_PATHWAY,PID\_EPHA2\_FWD\_PATHWAY  
GOBP\_FORMATION\_OF\_PRIMARY\_GERM\_LAYER,GOBP\_FORMATION\_OF\_PRIMARY\_GERM\_LAYER  
VERRECCHIA\_RESPONSE\_TO\_TGFB1\_C5,VERRECCHIA\_RESPONSE\_TO\_TGFB1\_C5  
GOBP\_MITRAL\_VALVE\_DEVELOPMENT,GOBP\_MITRAL\_VALVE\_DEVELOPMENT  
HP\_DYSLEXIA,HP\_DYSLEXIA  
GOBP\_MESENCHYME\_MORPHOGENESIS,GOBP\_MESENCHYME\_MORPHOGENESIS  
MIR6801\_3P,MIR6801\_3P  
HP\_CENTRAL\_DIABETES\_INSIPIDUS,HP\_CENTRAL\_DIABETES\_INSIPIDUS  
REACTOME\_RET\_SIGNALING,REACTOME\_RET\_SIGNALING  
GOBP\_BRANCHING\_MORPHOGENESIS\_OF\_AN\_EPITHELIAL\_TUBE,GOBP\_BRANCHING\_MORPHOGENESIS\_OF\_AN\_EPITHELIAL\_TUBE  
GOBP\_POSITIVE\_REGULATION\_OF\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION,GOBP\_POSITIVE\_REGULATION\_OF\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION  
GOMF\_HORMONE\_BINDING,GOMF\_HORMONE\_BINDING  
GOBP\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION,GOBP\_PATHWAY\_RESTRICTED\_SMAD\_PROTEIN\_PHOSPHORYLATION  
CADWELL\_ATG16L1\_TARGETS\_UP,CADWELL\_ATG16L1\_TARGETS\_UP  
MYLLYKANGAS\_AMPLIFICATION\_HOT\_SPOT\_9,MYLLYKANGAS\_AMPLIFICATION\_HOT\_SPOT\_9  
GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_SUBSTRATE\_ADHESION,GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_SUBSTRATE\_ADHESION  
HP\_LIMITED\_ELBOW\_EXTENSION,HP\_LIMITED\_ELBOW\_EXTENSION  
WP\_BMP\_SIGNALING\_PATHWAY\_IN\_EYELID\_DEVELOPMENT,WP\_BMP\_SIGNALING\_PATHWAY\_IN\_EYELID\_DEVELOPMENT  
GOBP\_VENTRICULAR\_SEPTUM\_MORPHOGENESIS,GOBP\_VENTRICULAR\_SEPTUM\_MORPHOGENESIS  
HP\_SPOKEN\_WORD\_RECOGNITION\_DEFICIT,HP\_SPOKEN\_WORD\_RECOGNITION\_DEFICIT  
HP\_PORTAL\_VEIN\_THROMBOSIS,HP\_PORTAL\_VEIN\_THROMBOSIS  
GOBP\_POSITIVE\_REGULATION\_OF\_EPIDERMAL\_CELL\_DIFFERENTIATION,GOBP\_POSITIVE\_REGULATION\_OF\_EPIDERMAL\_CELL\_DIFFERENTIATION  
HP\_GOWERS\_SIGN,HP\_GOWERS\_SIGN  
WP\_ROLE\_OF\_OSX\_AND\_MIRNAS\_IN\_TOOTH\_DEVELOPMENT,WP\_ROLE\_OF\_OSX\_AND\_MIRNAS\_IN\_TOOTH\_DEVELOPMENT  
WP\_SCAPHOCEPHALY,HP\_SCAPHOCEPHALY  
GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_LIGHTYELLOW\_UP,GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_LIGHTYELLOW\_UP  
HE\_PTFEN\_TARGETS\_UP,HE\_PTFEN\_TARGETS\_UP  
GOBP\_POSITIVE\_REGULATION\_OF\_CALCIIUM\_ION\_TRANSPORT\_INTO\_CYTOSOL,GOBP\_POSITIVE\_REGULATION\_OF\_CALCIIUM\_ION\_TRANSPORT\_INTO\_CYTOSOL  
FIGUEROA\_AML\_METHYLATION\_CLUSTER\_2\_DN,FIGUEROA\_AML\_METHYLATION\_CLUSTER\_2\_DN  
GOBP\_NEGATIVE\_REGULATION\_OF\_HORMONE\_METABOLIC\_PROCESS,GOBP\_NEGATIVE\_REGULATION\_OF\_HORMONE\_METABOLIC\_PROCESS  
POOLA\_INVASIVE\_BREAST\_CANCER\_DN,POOLA\_INVASIVE\_BREAST\_CANCER\_DN  
BOGNI\_TREATMENT\_RELATED\_MYELOID\_LEUKEMIA\_DN,BOGNI\_TREATMENT\_RELATED\_MYELOID\_LEUKEMIA\_DN  
DING\_LUNG\_CANCER\_BY\_MUTATION\_RATE,DING\_LUNG\_CANCER\_BY\_MUTATION\_RATE  
CLASPER\_LYMPHATIC\_VESSELS\_DURING\_METASTASIS\_DN,CLASPER\_LYMPHATIC\_VESSELS\_DURING\_METASTASIS\_DN  
GOMF\_PHOSPHOLIPASE\_BINDING,GOMF\_PHOSPHOLIPASE\_BINDING  
GOBP\_HINDLIMB\_MORPHOGENESIS,GOBP\_HINDLIMB\_MORPHOGENESIS  
GOMF\_PEPTIDE\_HORMONE\_BINDING,GOMF\_PEPTIDE\_HORMONE\_BINDING