

GSE2770\_TGFB\_AND\_IL4\_VS\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_TGFB\_AND\_IL4\_VS\_IL12\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP  
GSE22886\_DAY0\_VS\_DAY7\_MONOCYTE\_IN\_CULTURE\_UP, GSE22886\_DAY0\_VS\_DAY7\_MONOCYTE\_IN\_CULTURE\_UP  
GSE26559\_TCF1\_KO\_VS\_WT\_LIN\_NEG\_CELL\_UP, GSE26559\_TCF1\_KO\_VS\_WT\_LIN\_NEG\_CELL\_UP  
GSE29949\_CD8\_NEG\_DC\_SPLEEN\_VS\_MONOCYTE\_BONE\_MARROW\_UP, GSE29949\_CD8\_NEG\_DC\_SPLEEN\_VS\_MONOCYTE\_BONE\_MARROW\_UP  
GSE29949\_CD8\_POS\_DC\_SPLEEN\_VS\_DC\_BRAIN\_UP, GSE29949\_CD8\_POS\_DC\_SPLEEN\_VS\_DC\_BRAIN\_UP  
GSE29949\_DC\_BRAIN\_VS\_MONOCYTE\_BONE\_MARROW\_UP, GSE29949\_DC\_BRAIN\_VS\_MONOCYTE\_BONE\_MARROW\_UP  
GSE22601\_IMMATURE\_CD4\_SINGLE\_POSITIVE\_VS\_CD8\_SINGLE\_POSITIVE\_THYMOCYTE\_DN, GSE22601\_IMMATURE\_CD4\_SINGLE\_POSITIVE\_VS\_CD8\_SINGLE\_POSITIVE\_THYMOCYTE\_DN  
GSE22140\_GERMFREE\_VS\_SPF\_MOUSE\_CD4\_TCELL\_UP, GSE22140\_GERMFREE\_VS\_SPF\_MOUSE\_CD4\_TCELL\_UP  
GSE36888\_UNTREATED\_VS\_IL2\_TREATED\_TCELL\_6H\_UP, GSE36888\_UNTREATED\_VS\_IL2\_TREATED\_TCELL\_6H\_UP  
GSE21360\_PRIMARY\_VS\_TERTIARY\_MEMORY\_CD8\_TCELL\_DN, GSE21360\_PRIMARY\_VS\_TERTIARY\_MEMORY\_CD8\_TCELL\_DN  
GSE43955\_TH0\_VS\_TGFB\_IL6\_IL23\_THI7\_ACT\_CD4\_TCELL\_60H\_DN, GSE43955\_TH0\_VS\_TGFB\_IL6\_IL23\_THI7\_ACT\_CD4\_TCELL\_60H\_DN  
GSE23568\_CTRL\_TRANSDUCED\_VS\_WT\_CD8\_TCELL\_DN, GSE23568\_CTRL\_TRANSDUCED\_VS\_WT\_CD8\_TCELL\_DN  
GSE21670\_STAT3\_KO\_VS\_WT\_CD4\_TCELL\_DN, GSE21670\_STAT3\_KO\_VS\_WT\_CD4\_TCELL\_DN  
MODULE\_321, MODULE\_321  
BROCKE\_APOPTOSIS\_REVERSED\_BY\_IL6, BROCKE\_APOPTOSIS\_REVERSED\_BY\_IL6  
GSE360\_HIGH\_VS\_LOW\_DOSE\_B\_MALAYI\_DC\_UP, GSE360\_HIGH\_VS\_LOW\_DOSE\_B\_MALAYI\_DC\_UP  
GSE2770\_UNTREATED\_VS\_TGFB\_AND\_IL4\_TREATED\_ACT\_CD4\_TCELL\_48H\_DN, GSE2770\_UNTREATED\_VS\_TGFB\_AND\_IL4\_TREATED\_ACT\_CD4\_TCELL\_48H\_DN  
GSE43955\_TH0\_VS\_TGFB\_IL6\_THI7\_ACT\_CD4\_TCELL\_4H\_UP, GSE43955\_TH0\_VS\_TGFB\_IL6\_THI7\_ACT\_CD4\_TCELL\_4H\_UP  
GSE26030\_UNSTIM\_VS\_RESTIM\_THI7\_DAY5\_POST\_POLARIZATION\_DN, GSE26030\_UNSTIM\_VS\_RESTIM\_THI7\_DAY5\_POST\_POLARIZATION\_DN  
GSE360\_LOW\_DOSE\_B\_MALAYI\_VS\_M\_TUBERCULOSIS\_MAC\_DN, GSE360\_LOW\_DOSE\_B\_MALAYI\_VS\_M\_TUBERCULOSIS\_MAC\_DN  
GSE29949\_CD8\_POS\_DC\_SPLEEN\_VS\_MONOCYTE\_BONE\_MARROW\_DN, GSE29949\_CD8\_POS\_DC\_SPLEEN\_VS\_MONOCYTE\_BONE\_MARROW\_DN  
AIZARANI\_LIVER\_C21\_STELLATE\_CELLS\_1, AIZARANI\_LIVER\_C21\_STELLATE\_CELLS\_1  
GSE17974\_IL4\_AND\_ANTI\_IL12\_VS\_UNTREATED\_2H\_ACT\_CD4\_TCELL\_UP, GSE17974\_IL4\_AND\_ANTI\_IL12\_VS\_UNTREATED\_2H\_ACT\_CD4\_TCELL\_UP  
AIZARANI\_LIVER\_C39\_EPCAM\_POS\_BILE\_DUCT\_CELLS\_4, AIZARANI\_LIVER\_C39\_EPCAM\_POS\_BILE\_DUCT\_CELLS\_4  
PDGF\_UP.V1\_UP, PDGF\_UP.V1\_UP  
MIR224\_5P, MIR224\_5P  
GSE1460\_CORD\_VS\_ADULT\_BLOOD\_NAIVE\_CD4\_TCELL\_UP, GSE1460\_CORD\_VS\_ADULT\_BLOOD\_NAIVE\_CD4\_TCELL\_UP  
GSE13306\_RA\_VS\_UNTREATED\_MEM\_CD4\_TCELL\_DN, GSE13306\_RA\_VS\_UNTREATED\_MEM\_CD4\_TCELL\_DN  
GSE22025\_UNTREATED\_VS\_TGFB1\_TREATED\_CD4\_TCELL\_UP, GSE22025\_UNTREATED\_VS\_TGFB1\_TREATED\_CD4\_TCELL\_UP  
TONKS\_TARGETS\_OF\_RUNX1\_RUNX1T1\_FUSION\_ERYTHROCYTE\_UP, TONKS\_TARGETS\_OF\_RUNX1\_RUNX1T1\_FUSION\_ERYTHROCYTE\_UP  
GSE3982\_EOSINOPHIL\_VS\_DC\_UP, GSE3982\_EOSINOPHIL\_VS\_DC\_UP  
GSE22025\_PROGESTERONE\_VS\_TGFB1\_AND\_PROGESTERONE\_TREATED\_CD4\_TCELL\_DN, GSE22025\_PROGESTERONE\_VS\_TGFB1\_AND\_PROGESTERONE\_TREATED\_CD4\_TCELL\_DN  
CHIANG\_LIVER\_CANCER\_SUBCLASS\_CTNNB1\_DN, CHIANG\_LIVER\_CANCER\_SUBCLASS\_CTNNB1\_DN  
GSE1925\_CTRL\_VS\_3H\_IFNG\_STIM\_IFNG\_PRIMED\_MACROPHAGE\_DN, GSE1925\_CTRL\_VS\_3H\_IFNG\_STIM\_IFNG\_PRIMED\_MACROPHAGE\_DN  
GSE29949\_CD8\_NEG\_DC\_SPLEEN\_VS\_CD8\_POS\_DC\_SPLEEN\_DN, GSE29949\_CD8\_NEG\_DC\_SPLEEN\_VS\_CD8\_POS\_DC\_SPLEEN\_DN  
BURTON\_ADIPOGENESIS\_8, BURTON\_ADIPOGENESIS\_8  
AGCATT\_A\_MIR155, AGCATT\_A\_MIR155  
GSE17301\_IFNA2\_VS\_IFNA2\_AND\_ACD3\_ACD28\_STIM\_CD8\_TCELL\_DN, GSE17301\_IFNA2\_VS\_IFNA2\_AND\_ACD3\_ACD28\_STIM\_CD8\_TCELL\_DN  
GSE19941\_UNSTIM\_VS\_LPS\_AND\_IL10\_STIM\_IL10\_KO\_MACROPHAGE\_UP, GSE19941\_UNSTIM\_VS\_LPS\_AND\_IL10\_STIM\_IL10\_KO\_MACROPHAGE\_UP  
AIZARANI\_LIVER\_C24\_EPCAM\_POS\_BILE\_DUCT\_CELLS\_3, AIZARANI\_LIVER\_C24\_EPCAM\_POS\_BILE\_DUCT\_CELLS\_3  
WANG\_ESOPHAGUS\_CANCER\_VS\_NORMAL\_DN, WANG\_ESOPHAGUS\_CANCER\_VS\_NORMAL\_DN  
MODULE\_206, MODULE\_206  
GSE30153\_LUPUS\_VS\_HEALTHY\_DONOR\_BCELL\_DN, GSE30153\_LUPUS\_VS\_HEALTHY\_DONOR\_BCELL\_DN  
PID\_CXCR4\_PATHWAY, PID\_CXCR4\_PATHWAY  
HELLER\_SILENCED\_BY\_METHYLATION\_DN, HELLER\_SILENCED\_BY\_METHYLATION\_DN  
P53\_DN.V1\_DN, P53\_DN.V1\_DN  
GSE33374\_CD8\_ALPHAALPHA\_VS\_ALPHABETA\_CD161\_HIGH\_TCELL\_UP, GSE33374\_CD8\_ALPHAALPHA\_VS\_ALPHABETA\_CD161\_HIGH\_TCELL\_UP  
HP\_JAUNDICE, HP\_JAUNDICE  
ACOSTA\_PROLIFERATION\_INDEPENDENT\_MYC\_TARGETS\_DN, ACOSTA\_PROLIFERATION\_INDEPENDENT\_MYC\_TARGETS\_DN  
WP\_ADIPOGENESIS, WP\_ADIPOGENESIS  
DURANTE\_ADULT\_OLFACTORY\_NEUROEPITHELIUM\_VASCULAR\_SMOOTH\_MUSCLE\_CELLS, DURANTE\_ADULT\_OLFACTORY\_NEUROEPITHELIUM\_VASCULAR\_SMOOTH\_MUSCLE\_CELLS  
GSE19941\_UNSTIM\_VS\_LPS\_STIM\_IL10\_KO\_MACROPHAGE\_UP, GSE19941\_UNSTIM\_VS\_LPS\_STIM\_IL10\_KO\_MACROPHAGE\_UP  
MIR5590\_5P, MIR5590\_5P  
GSE34156\_UNTREATED\_VS\_24H\_NOD2\_AND\_TLR1\_TLR2\_LIGAND\_TREATED\_MONOCYTE\_UP, GSE34156\_UNTREATED\_VS\_24H\_NOD2\_AND\_TLR1\_TLR2\_LIGAND\_TREATED\_MONOCYTE\_UP  
KEGG\_AXON\_GUIDANCE, KEGG\_AXON\_GUIDANCE  
REACTOME\_SIGNALING\_BY\_MET, REACTOME\_SIGNALING\_BY\_MET  
EBAUER\_TARGETS\_OF\_PAX3\_FOXO1\_FUSION\_UP, EBAUER\_TARGETS\_OF\_PAX3\_FOXO1\_FUSION\_UP  
FRASOR\_RESPONSE\_TO ESTRADIOL\_DN, FRASOR\_RESPONSE\_TO ESTRADIOL\_DN  
REACTOME\_RESPONSE\_TO\_ELEVATED\_PLATELET\_CYTOSOLIC\_CA2, REACTOME\_RESPONSE\_TO\_ELEVATED\_PLATELET\_CYTOSOLIC\_CA2  
HP\_METAPHYSEAL\_WIDENING, HP\_METAPHYSEAL\_WIDENING  
REACTOME\_SEMAPHORIN\_INTERACTIONS, REACTOME\_SEMAPHORIN\_INTERACTIONS  
AMIT\_EGF\_RESPONSE\_40\_HELA, AMIT\_EGF\_RESPONSE\_40\_HELA  
CHIARADONNA\_NEOPLASTIC\_TRANSFORMATION\_KRAS\_CDC25\_DN, CHIARADONNA\_NEOPLASTIC\_TRANSFORMATION\_KRAS\_CDC25\_DN  
GTAAAG\_MIR302B, GTAAAG\_MIR302B  
BMI1\_DN\_MEL18\_DN.V1\_DN, BMI1\_DN\_MEL18\_DN.V1\_DN  
GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CELL\_ADHESION, GOBP\_NEGATIVE\_REGULATION\_OF\_CELL\_CELL\_ADHESION  
GSE13229\_IMM\_VS\_INTMATURE\_NKCELL\_UP, GSE13229\_IMM\_VS\_INTMATURE\_NKCELL\_UP  
REACTOME\_SIGNALING\_BY\_PTK6, REACTOME\_SIGNALING\_BY\_PTK6  
MIR5579\_5P, MIR5579\_5P  
HP\_AMBLYOPIA, HP\_AMBLYOPIA  
SREBP1\_02, SREBP1\_02  
GOBP\_NEGATIVE\_REGULATION\_OF\_LYMPHOCYTE\_ACTIVATION, GOBP\_NEGATIVE\_REGULATION\_OF\_LYMPHOCYTE\_ACTIVATION  
GSE29949\_CD8\_NEG\_DC\_SPLEEN\_VS\_CD8\_POS\_DC\_SPLEEN\_UP, GSE29949\_CD8\_NEG\_DC\_SPLEEN\_VS\_CD8\_POS\_DC\_SPLEEN\_UP  
HP ABDOMINAL\_DISTENTION, HP ABDOMINAL\_DISTENTION  
GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_MAGENTA\_UP, GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_MAGENTA\_UP  
HP\_ANOPHTHALMIA, HP\_ANOPHTHALMIA  
GOBP\_TISSUE\_REMODELING, GOBP\_TISSUE\_REMODELING  
GSE6269\_HEALTHY\_VS\_E\_COLI\_INF\_PBMCDN, GSE6269\_HEALTHY\_VS\_E\_COLI\_INF\_PBMCDN  
BOSCO\_TH1\_CYTOTOXIC\_MODULE, BOSCO\_TH1\_CYTOTOXIC\_MODULE  
GOCC\_SPECIFIC\_GRANULE\_LUMEN, GOCC\_SPECIFIC\_GRANULE\_LUMEN  
GOBP\_ORGAN\_GROWTH, GOBP\_ORGAN\_GROWTH  
MODULE\_85, MODULE\_85  
MODULE\_259, MODULE\_259  
KYNG\_RESPONSE\_TO\_H2O2\_VIA\_ERCC6\_UP, KYNG\_RESPONSE\_TO\_H2O2\_VIA\_ERCC6\_UP  
KONDO\_PROSTATE\_CANCER\_HCP\_WITH\_H3K27ME3, KONDO\_PROSTATE\_CANCER\_HCP\_WITH\_H3K27ME3  
REACTOME\_PERK\_REGULATES\_GENE\_EXPRESSION, REACTOME\_PERK\_REGULATES\_GENE\_EXPRESSION  
GOCC\_CLUSTER\_OF\_ACTIN\_BASED\_CELL\_PROJECTIONS, GOCC\_CLUSTER\_OF\_ACTIN\_BASED\_CELL\_PROJECTIONS  
GOBP\_MALE\_SEX\_DIFFERENTIATION, GOBP\_MALE\_SEX\_DIFFERENTIATION  
HALMOS\_CEBPA\_TARGETS\_UP, HALMOS\_CEBPA\_TARGETS\_UP  
GOBP\_ACETYL\_COA\_METABOLIC\_PROCESS, GOBP\_ACETYL\_COA\_METABOLIC\_PROCESS  
MIR3665, MIR3665  
HP\_WORMIAN\_BONES, HP\_WORMIAN\_BONES  
BUSSLINGER\_GASTRIC\_G\_CELLS, BUSSLINGER\_GASTRIC\_G\_CELLS  
REACTOME\_CD28\_CO\_STIMULATION, REACTOME\_CD28\_CO\_STIMULATION  
GOBP\_MATERNAL\_PLACENTA\_DEVELOPMENT, GOBP\_MATERNAL\_PLACENTA\_DEVELOPMENT  
REACTOME\_CLASS\_1\_PEROXISOMAL\_MEMBRANE\_PROTEIN\_IMPORT, REACTOME\_CLASS\_1\_PEROXISOMAL\_MEMBRANE\_PROTEIN\_IMPORT  
HP\_LOWER\_LIMB\_AMYOTROPHY, HP\_LOWER\_LIMB\_AMYOTROPHY  
GOCC\_BRUSH\_BORDER, GOCC\_BRUSH\_BORDER  
GOBP\_BONE\_REMODELING, GOBP\_BONE\_REMODELING  
MODULE\_488, MODULE\_488  
MIR376A\_5P, MIR376A\_5P  
GOBP\_DECIDUALIZATION, GOBP\_DECIDUALIZATION  
XU\_RESPONSE\_TO\_TRETINOIN\_AND\_NSC682994\_UP, XU\_RESPONSE\_TO\_TRETINOIN\_AND\_NSC682994\_UP  
REACTOME\_MET\_PROMOTES\_CELL\_MOTILITY, REACTOME\_MET\_PROMOTES\_CELL\_MOTILITY  
GOCC\_PROTEIN\_COMPLEX\_INVOLVED\_IN\_CELL\_ADHESION, GOCC\_PROTEIN\_COMPLEX\_INVOLVED\_IN\_CELL\_ADHESION  
REACTOME\_CELL\_EXTRACELLULAR\_MATRIX\_INTERACTIONS, REACTOME\_CELL\_EXTRACELLULAR\_MATRIX\_INTERACTIONS  
WP\_NUCLEAR\_RECEPTORS\_IN\_LIPID\_METABOLISM\_AND\_TOXICITY, WP\_NUCLEAR\_RECEPTORS\_IN\_LIPID\_METABOLISM\_AND\_TOXICITY  
JECHLINGER\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_DN, JECHLINGER\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_DN  
BIOCARTA\_NUCLEARRS\_PATHWAY, BIOCARTA\_NUCLEARRS\_PATHWAY  
HP\_SPEECH\_ARTICULATION\_DIFFICULTIES, HP\_SPEECH\_ARTICULATION\_DIFFICULTIES  
SABATES\_COLORECTAL\_ADENOMA\_UP, SABATES\_COLORECTAL\_ADENOMA\_UP  
GOMF\_CERAMIDE\_BINDING, GOMF\_CERAMIDE\_BINDING  
GOBP\_NEGATIVE\_REGULATION\_OF\_T\_CELL\_PROLIFERATION, GOBP\_NEGATIVE\_REGULATION\_OF\_T\_CELL\_PROLIFERATION  
HP\_DELAYED\_ERUPTION\_OF\_PRIMARY\_TEETH, HP\_DELAYED\_ERUPTION\_OF\_PRIMARY\_TEETH  
CUI\_DEVELOPING\_HEART\_CARDIAC\_FIBROBLASTS, CUI\_DEVELOPING\_HEART\_CARDIAC\_FIBROBLASTS  
GOCC\_CHITOSOME, GOCC\_CHITOSOME  
GOBP\_NEGATIVE\_REGULATION\_OF\_CALCIIUM\_MEDIATED\_SIGNALING, GOBP\_NEGATIVE\_REGULATION\_OF\_CALCIIUM\_MEDIATED\_SIGNALING  
GOBP\_SYNCYTIIUM\_FORMATION, GOBP\_SYNCYTIIUM\_FORMATION  
REACTOME\_RESPONSE\_OF\_EIF2AK1\_HRI\_TO\_HEME\_DEFICIENCY, REACTOME\_RESPONSE\_OF\_EIF2AK1\_HRI\_TO\_HEME\_DEFICIENCY  
HP\_HEMOPHAGOCYTOSIS, HP\_HEMOPHAGOCYTOSIS  
GOCC\_INTERSTITIAL\_MATRIX, GOCC\_INTERSTITIAL\_MATRIX  
BALLIF\_DEVELOPMENTAL\_DISABILITY\_P16\_P12\_DELETION, BALLIF\_DEVELOPMENTAL\_DISABILITY\_P16\_P12\_DELETION  
CDPCR3\_01, CDPCR3\_01  
GOBP\_ADENYLATE\_CYCLASE\_INHIBITING\_G\_PROTEIN\_COUPLED\_RECEPTOR\_SIGNALING\_PATHWAY, GOBP\_ADENYLATE\_CYCLASE\_INHIBITING\_G\_PROTEIN\_COUPLED\_RECEPTOR\_SIGNALING\_PATHWAY  
GOBP\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_DEPOLYMERIZATION, GOBP\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_DEPOLYMERIZATION  
GOBP\_COLLAGEN\_CATABOLIC\_PROCESS, GOBP\_COLLAGEN\_CATABOLIC\_PROCESS  
GOCC\_MULTIVESICULAR\_BODY\_MEMBRANE, GOCC\_MULTIVESICULAR\_BODY\_MEMBRANE  
REACTOME\_REGULATION\_OF\_PYRUVATE\_DEHYDROGENASE\_PDH\_COMPLEX, REACTOME\_REGULATION\_OF\_PYRUVATE\_DEHYDROGENASE\_PDH\_COMPLEX  
REACTOME\_RHO\_GTPASES\_ACTIVATE\_PAKS, REACTOME\_RHO\_GTPASES\_ACTIVATE\_PAKS  
LIANG\_HEMATOPOIESIS\_STEM\_CELL\_NUMBER\_QTL, LIANG\_HEMATOPOIESIS\_STEM\_CELL\_NUMBER\_QTL  
GNF2\_SERPINB5, GNF2\_SERPINB5  
GOBP\_IRON\_IION\_TRANSMEMBRANE\_TRANSPORT, GOBP\_IRON\_IION\_TRANSMEMBRANE\_TRANSPORT  
REACTOME\_THROMBOXANE\_SIGNALLING\_THROUGH\_TP\_RECEPTOR, REACTOME\_THROMBOXANE\_SIGNALLING\_THROUGH\_TP\_RECEPTOR  
MIR210\_5P, MIR210\_5P  
GOMF\_CYTOSKELETAL\_ANCHOR\_ACTIVITY, GOMF\_CYTOSKELETAL\_ANCHOR\_ACTIVITY  
GOBP\_REGULATION\_OF\_MUSCLE\_ORGAN\_DEVELOPMENT, GOBP\_REGULATION\_OF\_MUSCLE\_ORGAN\_DEVELOPMENT  
REACTOME\_HIGHLY\_CALCIIUM\_PERMEABLE\_NICOTINIC\_ACETYLCHOLINE\_RECEPTORS, REACTOME\_HIGHLY\_CALCIIUM\_PERMEABLE\_NICOTINIC\_ACETYLCHOLINE\_RECEPTORS  
HP\_VISUAL\_ACUITY\_TEST\_ABNORMALITY, HP\_VISUAL\_ACUITY\_TEST\_ABNORMALITY  
ALONSO\_METASTASIS\_NEURAL\_UP, ALONSO\_METASTASIS\_NEURAL\_UP  
CHEN\_ETV5\_TARGETS\_SERTOLI, CHEN\_ETV5\_TARGETS\_SERTOLI  
LIU\_VAV3\_PROSTATE\_CARCINOGENESIS\_DN, LIU\_VAV3\_PROSTATE\_CARCINOGENESIS\_DN  
HP\_ANODONTIA, HP\_ANODONTIA  
REACTOME\_ABC\_TRANSPORTERS\_IN\_LIPID\_HOMEOSTASIS, REACTOME\_ABC\_TRANSPORTERS\_IN\_LIPID\_HOMEOSTASIS  
HP\_RETINAL\_NEOVASCULARIZATION, HP\_RETINAL\_NEOVASCULARIZATION  
GOBP\_REGULATION\_OF\_MEMBRANE\_REPOLARIZATION\_DURING\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL, GOBP\_REGULATION\_OF\_MEMBRANE\_REPOLARIZATION\_DURING\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL  
REACTOME\_EPHRIN\_SIGNALING, REACTOME\_EPHRIN\_SIGNALING