

DIATE\_FILAMENT, GO\_INTERMEDIATE\_FILAMENT

GO\_KERATIN\_FILAMENT, GO\_KERATIN\_FILAMENT  
GO\_CELL\_JUNCTION\_ORGANIZATION, GO\_CELL\_JUNCTION\_ORGANIZATION  
KIM\_RESPONSE\_TO\_TSA\_AND\_DECITABINE\_UP, KIM\_RESPONSE\_TO\_TSA\_AND\_DECITABINE\_UP  
GO\_KERATINIZATION, GO\_KERATINIZATION  
ESC\_V6.5\_UP\_EARLY.V1\_DN, ESC\_V6.5\_UP\_EARLY.V1\_DN  
PID\_CASPASE\_PATHWAY, PID\_CASPASE\_PATHWAY  
LEF1\_UP.V1\_UP, LEF1\_UP.V1\_UP  
RICKMAN\_HEAD\_AND\_NECK\_CANCER\_E, RICKMAN\_HEAD\_AND\_NECK\_CANCER\_E  
chr21q22, chr21q22  
MODULE\_297, MODULE\_297  
chr17q12, chr17q12  
HINATA\_NFKB\_TARGETS\_KERATINOCYTE\_UP, HINATA\_NFKB\_TARGETS\_KERATINOCYTE\_UP  
MODULE\_357, MODULE\_357  
SENESE\_HDAC2\_TARGETS\_DN, SENESE\_HDAC2\_TARGETS\_DN  
MODULE\_154, MODULE\_154  
GSE41867\_DAY6\_VS\_DAY15\_LCMV\_CLONE13\_EFFECTOR\_CD8\_TCELL\_UP, GSE41867\_DAY6\_VS\_DAY15\_LCMV\_CLONE13\_EFFECTOR\_CD8\_TCELL\_UP  
GNF2\_SERPINB5, GNF2\_SERPINB5  
MODULE\_298, MODULE\_298  
BIOCARTA\_TNFR1\_PATHWAY, BIOCARTA\_TNFR1\_PATHWAY  
LE\_SKI\_TARGETS\_UP, LE\_SKI\_TARGETS\_UP  
MODULE\_157, MODULE\_157  
GO\_CELL\_CELL\_ADHERENS\_JUNCTION, GO\_CELL\_CELL\_ADHERENS\_JUNCTION  
GSE22443\_NAIVE\_VS\_ACT\_AND\_IL2\_TREATED\_CD8\_TCELL\_DN, GSE22443\_NAIVE\_VS\_ACT\_AND\_IL2\_TREATED\_CD8\_TCELL\_DN  
GO\_KERATINOCYTE\_DIFFERENTIATION, GO\_KERATINOCYTE\_DIFFERENTIATION  
GSE37532\_WT\_VS\_PPARG\_KO\_LN\_TREG\_DN, GSE37532\_WT\_VS\_PPARG\_KO\_LN\_TREG\_DN  
GSE19374\_UNINF\_VS\_LISTERIA\_INFECTED\_MACROPHAGE\_UP, GSE19374\_UNINF\_VS\_LISTERIA\_INFECTED\_MACROPHAGE\_UP  
MODULE\_68, MODULE\_68  
GSE5589\_WT\_VS\_IL10\_KO\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_180MIN\_DN, GSE5589\_WT\_VS\_IL10\_KO\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_180MIN\_DN  
MODULE\_111, MODULE\_111  
WANG\_BARRETTES\_ESOPHAGUS\_AND\_ESOPHAGUS\_CANCER\_DN, WANG\_BARRETTES\_ESOPHAGUS\_AND\_ESOPHAGUS\_CANCER\_DN  
KRAS.LUNG\_UP.V1\_DN, KRAS.LUNG\_UP.V1\_DN  
GO\_CELL\_JUNCTION\_ASSEMBLY, GO\_CELL\_JUNCTION\_ASSEMBLY  
DOANE\_BREAST\_CANCER\_ESR1\_DN, DOANE\_BREAST\_CANCER\_ESR1\_DN  
KEGG\_ARRHYTHMOGENIC\_RIGHT\_VENTRICULAR\_CARDIOMYOPATHY\_ARVC, KEGG\_ARRHYTHMOGENIC\_RIGHT\_VENTRICULAR\_CARDIOMYOPATHY\_ARVC  
TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_LOBULAR\_NORMAL\_UP, TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_LOBULAR\_NORMAL\_UP  
GO\_HEPATOCYTE\_APOPTOTIC\_PROCESS, GO\_HEPATOCYTE\_APOPTOTIC\_PROCESS  
TURASHVILI\_BREAST\_DUCTAL\_CARCINOMA\_VS\_LOBULAR\_NORMAL\_DN, TURASHVILI\_BREAST\_DUCTAL\_CARCINOMA\_VS\_LOBULAR\_NORMAL\_DN  
REACTOME\_APOPTOTIC\_EXECUTION\_PHASE, REACTOME\_APOPTOTIC\_EXECUTION\_PHASE  
GO\_REGULATION\_OF\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL, GO\_REGULATION\_OF\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL  
GO\_HETEROTYPIC\_CELL\_CELL\_ADHESION, GO\_HETEROTYPIC\_CELL\_CELL\_ADHESION  
SNF5\_DN.V1\_DN, SNF5\_DN.V1\_DN  
GO\_MULTICELLULAR\_ORGANISMAL\_SIGNALING, GO\_MULTICELLULAR\_ORGANISMAL\_SIGNALING  
GSE7459\_UNTREATED\_VS\_IL6\_TREATED\_ACT\_CD4\_TCELL\_DN, GSE7459\_UNTREATED\_VS\_IL6\_TREATED\_ACT\_CD4\_TCELL\_DN  
GO\_MULTICELLULAR\_ORGANISMAL\_WATER\_HOMEOSTASIS, GO\_MULTICELLULAR\_ORGANISMAL\_WATER\_HOMEOSTASIS  
GO\_CELL\_CELL\_SIGNALING\_INVOLVED\_IN\_CARDIAC\_CONDUCTION, GO\_CELL\_CELL\_SIGNALING\_INVOLVED\_IN\_CARDIAC\_CONDUCTION  
GO\_REGULATION\_OF\_VENTRICULAR\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL, GO\_REGULATION\_OF\_VENTRICULAR\_CARDIAC\_MUSCLE\_CELL\_ACTION\_POTENTIAL  
POOLA\_INVASIVE\_BREAST\_CANCER\_DN, POOLA\_INVASIVE\_BREAST\_CANCER\_DN  
MUELLER\_METHYLATED\_IN\_GLIOBLASTOMA, MUELLER\_METHYLATED\_IN\_GLIOBLASTOMA  
GO\_CARDIAC\_MUSCLE\_CELL\_CONTRACTION, GO\_CARDIAC\_MUSCLE\_CELL\_CONTRACTION  
GO\_REGULATION\_OF\_ACTIN\_FILAMENT\_BASED\_MOVEMENT, GO\_REGULATION\_OF\_ACTIN\_FILAMENT\_BASED\_MOVEMENT  
GO\_ACTIN\_FILAMENT\_BASED\_MOVEMENT, GO\_ACTIN\_FILAMENT\_BASED\_MOVEMENT  
AP2REP\_01, AP2REP\_01  
ACEVEDO\_LIVER\_CANCER\_WITH\_H3K9ME3\_UP, ACEVEDO\_LIVER\_CANCER\_WITH\_H3K9ME3\_UP  
TURASHVILI\_BREAST\_DUCTAL\_CARCINOMA\_VS\_DUCTAL\_NORMAL\_DN, TURASHVILI\_BREAST\_DUCTAL\_CARCINOMA\_VS\_DUCTAL\_NORMAL\_DN  
MODULE\_153, MODULE\_153  
CAR\_TNFRSF25, CAR\_TNFRSF25  
TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_DUCTAL\_NORMAL\_DN, TURASHVILI\_BREAST\_LOBULAR\_CARCINOMA\_VS\_DUCTAL\_NORMAL\_DN  
GSE9988\_LPS\_VS\_LOW\_LPS\_MONOCYTE\_UP, GSE9988\_LPS\_VS\_LOW\_LPS\_MONOCYTE\_UP  
GO\_STRUCTURAL\_CONSTITUENT\_OF\_CYTOSKELETON, GO\_STRUCTURAL\_CONSTITUENT\_OF\_CYTOSKELETON  
GO\_STRUCTURAL\_CONSTITUENT\_OF\_EYE\_LENS, GO\_STRUCTURAL\_CONSTITUENT\_OF\_EYE\_LENS  
GO\_CELL\_SUBSTRATE\_JUNCTION\_ASSEMBLY, GO\_CELL\_SUBSTRATE\_JUNCTION\_ASSEMBLY  
HUPER\_BREAST\_BASAL\_VS\_LUMINAL\_UP, HUPER\_BREAST\_BASAL\_VS\_LUMINAL\_UP  
GSE18148\_CFBF\_KO\_VS\_WT\_TREG\_UP, GSE18148\_CFBF\_KO\_VS\_WT\_TREG\_UP  
GNF2\_SPRR1B, GNF2\_SPRR1B  
GO\_SCAFFOLD\_PROTEIN\_BINDING, GO\_SCAFFOLD\_PROTEIN\_BINDING  
GO\_GLYCOPROTEIN\_COMPLEX, GO\_GLYCOPROTEIN\_COMPLEX  
GO\_PROTEIN\_BINDING\_INVOLVED\_IN\_CELL\_ADHESION, GO\_PROTEIN\_BINDING\_INVOLVED\_IN\_CELL\_ADHESION  
LIN\_SILENCED\_BY\_TUMOR\_MICROENVIRONMENT, LIN\_SILENCED\_BY\_TUMOR\_MICROENVIRONMENT  
GO\_CARDIAC\_CONDUCTION, GO\_CARDIAC\_CONDUCTION  
GO\_REGULATION\_OF\_HEART\_RATE, GO\_REGULATION\_OF\_HEART\_RATE  
GO\_INTERCALATED\_DISC, GO\_INTERCALATED\_DISC  
JI\_CARCINOGENESIS\_BY\_KRAS\_AND\_STK11\_UP, JI\_CARCINOGENESIS\_BY\_KRAS\_AND\_STK11\_UP  
GO\_CELL\_CELL\_CONTACT\_ZONE, GO\_CELL\_CELL\_CONTACT\_ZONE  
GSE10325\_MYELOID\_VS\_LUPUS\_MYELOID\_UP, GSE10325\_MYELOID\_VS\_LUPUS\_MYELOID\_UP  
GSE13887\_RESTING\_VS\_NO\_TREATED\_CD4\_TCELL\_UP, GSE13887\_RESTING\_VS\_NO\_TREATED\_CD4\_TCELL\_UP  
GU\_PDEF\_TARGETS\_DN, GU\_PDEF\_TARGETS\_DN