

KO\_MACROPHAGE\_2H\_LPS\_STIM\_UP, GSE30971\_WBP7\_HET\_VS\_KO\_MACROPHAGE\_2H\_LPS\_STIM\_UP

GSE21360\_TERTIARY\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_UP, GSE21360\_TERTIARY\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_UP  
GSE13306\_RA\_VS\_UNTREATED\_MEM\_CD4\_TCELL\_UP, GSE13306\_RA\_VS\_UNTREATED\_MEM\_CD4\_TCELL\_UP  
GSE20727\_CTRL\_VS\_ROS\_INH\_AND\_DNFB\_ALLERGEN\_TREATED\_DC\_UP, GSE20727\_CTRL\_VS\_ROS\_INH\_AND\_DNFB\_ALLERGEN\_TREATED\_DC\_UP  
GSE3982\_MAC\_VS\_BCELL\_UP, GSE3982\_MAC\_VS\_BCELL\_UP  
KRAS.DF.V1\_UP, KRAS.DF.V1\_UP  
BURTON\_ADIPOGENESIS\_9, BURTON\_ADIPOGENESIS\_9  
NABA\_ECM\_AFFILIATED, NABA\_ECM\_AFFILIATED  
GSE44732\_UNSTIM\_VS\_IL27\_STIM\_IMATURE\_DC\_UP, GSE44732\_UNSTIM\_VS\_IL27\_STIM\_IMATURE\_DC\_UP  
TGFB\_UP.V1\_DN, TGFB\_UP.V1\_DN  
GSE17721\_CTRL\_VS\_GARDIQUIMOD\_2H\_BMDC\_DN, GSE17721\_CTRL\_VS\_GARDIQUIMOD\_2H\_BMDC\_DN  
GSE20500\_RETINOIC\_ACID\_VS\_RARA\_ANTAGONIST\_TREATED\_CD4\_TCELL\_UP, GSE20500\_RETINOIC\_ACID\_VS\_RARA\_ANTAGONIST\_TREATED\_CD4\_TCELL\_UP  
CERVERA\_SDHB\_TARGETS\_2, CERVERA\_SDHB\_TARGETS\_2  
PTEN\_DN.V1\_UP, PTEN\_DN.V1\_UP  
MODULE\_180, MODULE\_180  
GSE39820\_IL1B\_IL6\_VS\_IL1B\_IL6\_IL23A\_TREATED\_CD4\_TCELL\_UP, GSE39820\_IL1B\_IL6\_VS\_IL1B\_IL6\_IL23A\_TREATED\_CD4\_TCELL\_UP  
GO\_NEGATIVE\_REGULATION\_OF\_CELL\_GROWTH, GO\_NEGATIVE\_REGULATION\_OF\_CELL\_GROWTH  
GO\_RESPONSE\_TO\_PROGESTERONE, GO\_RESPONSE\_TO\_PROGESTERONE  
MODULE\_96, MODULE\_96  
GO\_HUMORAL\_IMMUNE\_RESPONSE, GO\_HUMORAL\_IMMUNE\_RESPONSE  
MODULE\_410, MODULE\_410  
NEWMAN\_ERCC6\_TARGETS\_UP, NEWMAN\_ERCC6\_TARGETS\_UP  
GO\_ENDODERMAL\_CELL\_DIFFERENTIATION, GO\_ENDODERMAL\_CELL\_DIFFERENTIATION  
GO\_REGULATION\_OF\_BONE\_RESORPTION, GO\_REGULATION\_OF\_BONE\_RESORPTION  
ENGELMANN\_CANCER\_PROGENITORS\_DN, ENGELMANN\_CANCER\_PROGENITORS\_DN  
GO\_REGULATION\_OF\_PHOSPHOLIPASE\_ACTIVITY, GO\_REGULATION\_OF\_PHOSPHOLIPASE\_ACTIVITY  
GO\_ION\_ANTIPORTER\_ACTIVITY, GO\_ION\_ANTIPORTER\_ACTIVITY  
GO\_RESPIRATORY\_SYSTEM\_DEVELOPMENT, GO\_RESPIRATORY\_SYSTEM\_DEVELOPMENT  
MODULE\_495, MODULE\_495  
GO\_ENDODERM\_FORMATION, GO\_ENDODERM\_FORMATION  
GSE21927\_SPLEEN\_VS\_C26GM\_TUMOR\_MONOCYTE\_BALBC\_UP, GSE21927\_SPLEEN\_VS\_C26GM\_TUMOR\_MONOCYTE\_BALBC\_UP  
VALK\_AML\_CLUSTER\_12, VALK\_AML\_CLUSTER\_12  
GO\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_MATURATION, GO\_NEGATIVE\_REGULATION\_OF\_PROTEIN\_MATURATION  
GO\_RESPONSE\_TO\_GRAVITY, GO\_RESPONSE\_TO\_GRAVITY  
GSE40274\_EOS\_VS\_FOXP3\_AND\_EOS\_TRANSDUCE\_ACTIVATED\_CD4\_TCELL\_DN, GSE40274\_EOS\_VS\_FOXP3\_AND\_EOS\_TRANSDUCE\_ACTIVATED\_CD4\_TCELL\_DN  
WANG\_BARRETTES\_ESOPHAGUS\_UP, WANG\_BARRETTES\_ESOPHAGUS\_UP  
GO\_STEM\_CELL\_DIFFERENTIATION, GO\_STEM\_CELL\_DIFFERENTIATION  
HERNANDEZ\_ABERRANT\_MITOSIS\_BY\_DOCETACEL\_2NM\_DN, HERNANDEZ\_ABERRANT\_MITOSIS\_BY\_DOCETACEL\_2NM\_DN  
REACTOME\_INTRINSIC\_PATHWAY, REACTOME\_INTRINSIC\_PATHWAY  
KRISHNAN\_FURIN\_TARGETS\_UP, KRISHNAN\_FURIN\_TARGETS\_UP  
GO\_LOCOMOTORY\_EXPLORATION\_BEHAVIOR, GO\_LOCOMOTORY\_EXPLORATION\_BEHAVIOR  
GO\_BLOOD\_COAGULATION\_INTRINSIC\_PATHWAY, GO\_BLOOD\_COAGULATION\_INTRINSIC\_PATHWAY