

ULTIPOTENT\_VS\_PRO\_BCELL\_DN, GSE15330\_LYMPHOID\_MULTIPOTENT\_VS\_PRO\_BCELL\_DN

GSE36891\_UNSTIM\_VS\_POLYIC\_TLR3\_STIM\_PERITONEAL\_MACROPHAGE\_DN, GSE36891\_UNSTIM\_VS\_POLYIC\_TLR3\_STIM\_PERITONEAL\_MACROPHAGE\_DN  
GSE17721\_CTRL\_VS\_GARDIQUIMOD\_2H\_BMDC\_DN, GSE17721\_CTRL\_VS\_GARDIQUIMOD\_2H\_BMDC\_DN  
GSE30971\_WBP7\_HET\_VS\_KO\_MACROPHAGE\_2H\_LPS\_STIM\_DN, GSE30971\_WBP7\_HET\_VS\_KO\_MACROPHAGE\_2H\_LPS\_STIM\_DN  
GSE26669\_CD4\_VS\_CD8\_TCELL\_IN\_MLR\_UP, GSE26669\_CD4\_VS\_CD8\_TCELL\_IN\_MLR\_UP  
GSE17721\_LPS\_VS\_PAM3CSK4\_12H\_BMDC\_UP, GSE17721\_LPS\_VS\_PAM3CSK4\_12H\_BMDC\_UP  
GSE17721\_0.5H\_VS\_12H\_LPS\_BMDC\_DN, GSE17721\_0.5H\_VS\_12H\_LPS\_BMDC\_DN  
GSE23308\_CTRL\_VS\_CORTICOSTERONE\_TREATED\_MACROPHAGE\_DN, GSE23308\_CTRL\_VS\_CORTICOSTERONE\_TREATED\_MACROPHAGE\_DN  
GSE21063\_CTRL\_VS\_ANTI\_IGM\_STIM\_BCELL\_NFATC1\_KO\_3H\_UP, GSE21063\_CTRL\_VS\_ANTI\_IGM\_STIM\_BCELL\_NFATC1\_KO\_3H\_UP  
GSE1925\_CTRL\_VS\_24H\_IFNG\_STIM\_MACROPHAGE\_DN, GSE1925\_CTRL\_VS\_24H\_IFNG\_STIM\_MACROPHAGE\_DN  
GSE30971\_WBP7\_HET\_VS\_KO\_MACROPHAGE\_DN, GSE30971\_WBP7\_HET\_VS\_KO\_MACROPHAGE\_DN  
GSE46606\_IRF4HIGH\_VS\_IRF4MID\_CD40L\_IL2\_IL5\_DAY3\_STIMULATED\_BCELL\_DN, GSE46606\_IRF4HIGH\_VS\_IRF4MID\_CD40L\_IL2\_IL5\_DAY3\_STIMULATED\_BCELL\_DN  
GSE22935\_UNSTIM\_VS\_24H\_MBOVIS\_BCG\_STIM\_MACROPHAGE\_DN, GSE22935\_UNSTIM\_VS\_24H\_MBOVIS\_BCG\_STIM\_MACROPHAGE\_DN  
GSE32423\_CTRL\_VS\_IL4\_MEMORY\_CD8\_TCELL\_DN, GSE32423\_CTRL\_VS\_IL4\_MEMORY\_CD8\_TCELL\_DN  
MIR6715B\_3P, MIR6715B\_3P  
MIR488\_5P, MIR488\_5P  
GSE2585\_CTEC\_VS\_THYMIC\_DC\_DN, GSE2585\_CTEC\_VS\_THYMIC\_DC\_DN  
CTCNANGTGNY\_UNKNOWN, CTCNANGTGNY\_UNKNOWN  
GSE13229\_IMM\_VS\_INTMATURE\_NKCELL\_UP, GSE13229\_IMM\_VS\_INTMATURE\_NKCELL\_UP  
GSE46606\_DAY1\_VS\_DAY3\_CD40L\_IL2\_IL5\_STIMULATED\_BCELL\_UP, GSE46606\_DAY1\_VS\_DAY3\_CD40L\_IL2\_IL5\_STIMULATED\_BCELL\_UP  
HP\_ABNORMAL\_ENDOCARDIUM\_MORPHOLOGY, HP\_ABNORMAL\_ENDOCARDIUM\_MORPHOLOGY  
GOBP\_REGULATION\_OF\_CATION\_CHANNEL\_ACTIVITY, GOBP\_REGULATION\_OF\_CATION\_CHANNEL\_ACTIVITY  
VALK\_AML\_CLUSTER\_11, VALK\_AML\_CLUSTER\_11  
GOBP\_INORGANIC\_ANION\_TRANSMEMBRANE\_TRANSPORT, GOBP\_INORGANIC\_ANION\_TRANSMEMBRANE\_TRANSPORT  
GOBP\_MEMORY, GOBP\_MEMORY  
GOMF\_CHLORIDE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY, GOMF\_CHLORIDE\_TRANSMEMBRANE\_TRANSPORTER\_ACTIVITY  
HP\_BASAL\_GANGLIA\_CALCIFICATION, HP\_BASAL\_GANGLIA\_CALCIFICATION  
GSE18281\_PERIMEDULLARY\_CORTICAL\_REGION\_VS\_WHOLE\_MEDULLA\_THYMUS\_DN, GSE18281\_PERIMEDULLARY\_CORTICAL\_REGION\_VS\_WHOLE\_MEDULLA\_THYMUS\_DN  
GOCC\_CHLORIDE\_CHANNEL\_COMPLEX, GOCC\_CHLORIDE\_CHANNEL\_COMPLEX  
GOBP\_CHLORIDE\_TRANSPORT, GOBP\_CHLORIDE\_TRANSPORT  
ZHONG\_PFC\_C2\_SOX5\_BCL11B\_POS\_EXCITATORY\_NEURON, ZHONG\_PFC\_C2\_SOX5\_BCL11B\_POS\_EXCITATORY\_NEURON  
GOMF\_SERINE\_TYPE\_ENDOPEPTIDASE\_INHIBITOR\_ACTIVITY, GOMF\_SERINE\_TYPE\_ENDOPEPTIDASE\_INHIBITOR\_ACTIVITY  
GOBP\_POLY\_N\_ACETYLLACTOSAMINE\_METABOLIC\_PROCESS, GOBP\_POLY\_N\_ACETYLLACTOSAMINE\_METABOLIC\_PROCESS  
GOMF\_N\_ACETYLLACTOSAMINIDE\_BETA\_1\_3\_N\_ACETYLGLUCOSAMINYLTRANSFERASE\_ACTIVITY, GOMF\_N\_ACETYLLACTOSAMINIDE\_BETA\_1\_3\_N\_ACETYLGLUCOSAMINYLTRANSFERASE\_ACTIVITY  
GOBP\_RESPONSE\_TO\_INTERLEUKIN\_17, GOBP\_RESPONSE\_TO\_INTERLEUKIN\_17