

\_NOD\_FOXP3\_FUSION\_GFP\_TREG\_DN, GSE37605\_C57BL6\_VS\_NOD\_FOXP3\_FUSION\_GFP\_TREG\_DN

GSE46606\_UNSTIM\_VS\_CD40L\_IL2\_IL5\_1DAY\_STIMULATED\_IRF4MID\_SORTED\_BCELL\_UP, GSE46606\_UNSTIM\_VS\_CD40L\_IL2\_IL5\_1DAY\_STIMULATED\_IRF4MID\_SORTED\_BCELL\_UP  
HALLMARK\_EPITHELIAL\_MESENCHYMAL\_TRANSITION, HALLMARK\_EPITHELIAL\_MESENCHYMAL\_TRANSITION  
GSE37605\_TREG\_VS\_TCONV\_NOD\_FOXP3\_FUSION\_GFP\_DN, GSE37605\_TREG\_VS\_TCONV\_NOD\_FOXP3\_FUSION\_GFP\_DN  
GSE14769\_UNSTIM\_VS\_20MIN\_LPS\_BMDM\_DN, GSE14769\_UNSTIM\_VS\_20MIN\_LPS\_BMDM\_DN  
GSE22103\_UNSTIM\_VS\_LPS\_STIM\_NEUTROPHIL\_DN, GSE22103\_UNSTIM\_VS\_LPS\_STIM\_NEUTROPHIL\_DN  
GSE43955\_10H\_VS\_60H\_ACT\_CD4\_TCELL\_WITH\_TGFB\_IL6\_DN, GSE43955\_10H\_VS\_60H\_ACT\_CD4\_TCELL\_WITH\_TGFB\_IL6\_DN  
MTOR\_UP.V1\_DN, MTOR\_UP.V1\_DN  
GO\_NEPHRON\_DEVELOPMENT, GO\_NEPHRON\_DEVELOPMENT  
ESC\_J1\_UP\_EARLY.V1\_DN, ESC\_J1\_UP\_EARLY.V1\_DN  
GO\_MORPHOGENESIS\_OF\_A\_BRANCHING\_STRUCTURE, GO\_MORPHOGENESIS\_OF\_A\_BRANCHING\_STRUCTURE  
MODULE\_92, MODULE\_92  
GSE41176\_WT\_VS\_TAK1\_KO\_ANTI\_IGM\_STIM\_BCELL\_6H\_UP, GSE41176\_WT\_VS\_TAK1\_KO\_ANTI\_IGM\_STIM\_BCELL\_6H\_UP  
DALESSIO\_TSA\_RESPONSE, DALESSIO\_TSA\_RESPONSE  
WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_DN, WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_DN  
GO\_PHOTOPERIODISM, GO\_PHOTOPERIODISM  
WIEDERSCHAIN\_TARGETS\_OF\_BMI1\_AND\_PCGF2, WIEDERSCHAIN\_TARGETS\_OF\_BMI1\_AND\_PCGF2  
GSE36476\_YOUNG\_VS\_OLD\_DONOR\_MEMORY\_CD4\_TCELL\_DN, GSE36476\_YOUNG\_VS\_OLD\_DONOR\_MEMORY\_CD4\_TCELL\_DN  
BERENJENO\_ROCK\_SIGNALING\_NOT\_VIA\_RHOA\_UP, BERENJENO\_ROCK\_SIGNALING\_NOT\_VIA\_RHOA\_UP  
GO\_FORMATION\_OF\_PRIMARY\_GERM\_LAYER, GO\_FORMATION\_OF\_PRIMARY\_GERM\_LAYER  
GO\_KIDNEY\_EPITHELIUM\_DEVELOPMENT, GO\_KIDNEY\_EPITHELIUM\_DEVELOPMENT  
GO\_NEGATIVE\_REGULATION\_OF\_CHEMOTAXIS, GO\_NEGATIVE\_REGULATION\_OF\_CHEMOTAXIS  
GO\_NEGATIVE\_REGULATION\_OF\_GTPASE\_ACTIVITY, GO\_NEGATIVE\_REGULATION\_OF\_GTPASE\_ACTIVITY  
GO\_REGULATION\_OF\_CYCLIC\_NUCLEOTIDE\_METABOLIC\_PROCESS, GO\_REGULATION\_OF\_CYCLIC\_NUCLEOTIDE\_METABOLIC\_PROCESS  
GSE29164\_UNTREATED\_VS\_CD8\_TCELL\_TREATED\_MELANOMA\_DAY7\_DN, GSE29164\_UNTREATED\_VS\_CD8\_TCELL\_TREATED\_MELANOMA\_DAY7\_DN  
GO\_REGULATION\_OF\_CAMP\_METABOLIC\_PROCESS, GO\_REGULATION\_OF\_CAMP\_METABOLIC\_PROCESS  
GO\_PEPTIDE\_RECEPTOR\_ACTIVITY, GO\_PEPTIDE\_RECEPTOR\_ACTIVITY  
GO\_EAR\_DEVELOPMENT, GO\_EAR\_DEVELOPMENT  
SCHLESINGER\_METHYLATED\_DE\_NOVO\_IN\_CANCER, SCHLESINGER\_METHYLATED\_DE\_NOVO\_IN\_CANCER  
GO\_MODULATION\_OF\_GROWTH\_OF\_SYMBIONT\_INVOLVED\_IN\_INTERACTION\_WITH\_HOST, GO\_MODULATION\_OF\_GROWTH\_OF\_SYMBIONT\_INVOLVED\_IN\_INTERACTION\_WIT  
GSE3982\_BCELL\_VS\_NKCELL\_DN, GSE3982\_BCELL\_VS\_NKCELL\_DN  
GO\_RENAL\_TUBULE\_DEVELOPMENT, GO\_RENAL\_TUBULE\_DEVELOPMENT  
GO\_RETINA\_MORPHOGENESIS\_IN\_CAMERA\_TYPE\_EYE, GO\_RETINA\_MORPHOGENESIS\_IN\_CAMERA\_TYPE\_EYE  
YNTTTNNNANGCARM\_UNKNOWN, YNTTTNNNANGCARM\_UNKNOWN  
GO\_REGULATION\_OF\_INFLAMMATORY\_RESPONSE\_TO\_ANTIGENIC\_STIMULUS, GO\_REGULATION\_OF\_INFLAMMATORY\_RESPONSE\_TO\_ANTIGENIC\_STIMULUS  
GO\_EAR\_MORPHOGENESIS, GO\_EAR\_MORPHOGENESIS  
GO\_CHEMOKINE\_ACTIVITY, GO\_CHEMOKINE\_ACTIVITY  
GO\_REGULATION\_OF\_SYSTEMIC\_ARTERIAL\_BLOOD\_PRESSURE\_MEDIATED\_BY\_A\_CHEMICAL\_SIGNAL, GO\_REGULATION\_OF\_SYSTEMIC\_ARTERIAL\_BLOOD\_PRESSURE\_MEDIATE  
OXFORD\_RALB\_TARGETS\_UP, OXFORD\_RALB\_TARGETS\_UP  
GO\_LYMPHOCYTE\_CHEMOTAXIS, GO\_LYMPHOCYTE\_CHEMOTAXIS  
WANG\_NFKB\_TARGETS, WANG\_NFKB\_TARGETS  
GO\_POSITIVE\_REGULATION\_OF\_MUSCLE\_CONTRACTION, GO\_POSITIVE\_REGULATION\_OF\_MUSCLE\_CONTRACTION