

IL2\_KO\_CD4\_TCELL\_SCURFY\_MOUSE\_DN, GSE23398\_WT\_VS\_IL2\_KO\_CD4\_TCELL\_SCURFY\_MOUSE\_DN

GSE39820\_CTRL\_VS\_IL1B\_IL6\_IL23A\_CD4\_TCELL\_DN, GSE39820\_CTRL\_VS\_IL1B\_IL6\_IL23A\_CD4\_TCELL\_DN  
GSE25677\_R848\_VS\_MPL\_AND\_R848\_STIM\_BCELL\_DN, GSE25677\_R848\_VS\_MPL\_AND\_R848\_STIM\_BCELL\_DN  
GSE3039\_NKT\_CELL\_VS\_ALPHAALPHA\_CD8\_TCELL\_UP, GSE3039\_NKT\_CELL\_VS\_ALPHAALPHA\_CD8\_TCELL\_UP  
MIR194\_5P, MIR194\_5P  
GSE43955\_TGFB\_IL6\_VS\_TGFB\_IL6\_IL23\_TH17\_ACT\_CD4\_TCELL\_52H\_UP, GSE43955\_TGFB\_IL6\_VS\_TGFB\_IL6\_IL23\_TH17\_ACT\_CD4\_TCELL\_52H\_UP  
ESC\_V6.5\_UP\_LATE.V1\_UP, ESC\_V6.5\_UP\_LATE.V1\_UP  
MIR152\_5P, MIR152\_5P  
GSE24972\_WT\_VS\_IRF8\_KO\_MARGINAL\_ZONE\_SPLEEN\_BCELL\_DN, GSE24972\_WT\_VS\_IRF8\_KO\_MARGINAL\_ZONE\_SPLEEN\_BCELL\_DN  
MIR200C\_5P, MIR200C\_5P  
MIR627\_5P, MIR627\_5P  
HOEK\_MYELOID\_DENDRITIC\_CELL\_2011\_2012\_TIV\_ADULT\_1DY\_UP, HOEK\_MYELOID\_DENDRITIC\_CELL\_2011\_2012\_TIV\_ADULT\_1DY\_UP  
GOCC\_ATPASE\_DEPENDENT\_TRANSMEMBRANE\_TRANSPORT\_COMPLEX, GOCC\_ATPASE\_DEPENDENT\_TRANSMEMBRANE\_TRANSPORT\_COMPLEX  
GOCC\_SODIUM\_POTASSIUM\_EXCHANGING\_ATPASE\_COMPLEX, GOCC\_SODIUM\_POTASSIUM\_EXCHANGING\_ATPASE\_COMPLEX  
HUANG\_FOXA2\_TARGETS\_UP, HUANG\_FOXA2\_TARGETS\_UP  
GOBP\_NEURON\_RECOGNITION, GOBP\_NEURON\_RECOGNITION  
GOBP\_INTRACILIARY\_RETROGRADE\_TRANSPORT, GOBP\_INTRACILIARY\_RETROGRADE\_TRANSPORT  
chr5q11, chr5q11  
GOBP\_SODIUM\_ION\_EXPORT\_ACROSS\_PLASMA\_MEMBRANE, GOBP\_SODIUM\_ION\_EXPORT\_ACROSS\_PLASMA\_MEMBRANE  
HP\_ABNORMAL\_SHARPEY\_FIBER\_MORPHOLOGY, HP\_ABNORMAL\_SHARPEY\_FIBER\_MORPHOLOGY  
HP\_CHONDROCALCINOSIS, HP\_CHONDROCALCINOSIS