


**TLR1\_2\_STIM\_MONOCYTE\_3H\_DN, GSE8921\_UNSTIM\_VS\_TLR1\_2\_STIM\_MONOCYTE\_3H\_DN**



GSE24972\_WT\_VS\_IRF8\_KO\_MARGINAL\_ZONE\_SPLEEN\_BCELL\_UP, GSE24972\_WT\_VS\_IRF8\_KO\_MARGINAL\_ZONE\_SPLEEN\_BCELL\_UP  
GSE15330\_HSC\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_IKAROS\_KO\_UP, GSE15330\_HSC\_VS\_GRANULOCYTE\_MONOCYTE\_PROGENITOR\_IKAROS\_KO\_UP  
GSE28737\_WT\_VS\_BCL6\_HET\_MARGINAL\_ZONE\_BCELL\_UP, GSE28737\_WT\_VS\_BCL6\_HET\_MARGINAL\_ZONE\_BCELL\_UP  
GSE5542\_UNTREATED\_VS\_IFNA\_AND\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_6H\_UP, GSE5542\_UNTREATED\_VS\_IFNA\_AND\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_6H\_UP  
GSE21927\_SPLEEN\_MONOCYTE\_VS\_GMCSF\_GCSF\_BONE\_MARROW\_UP, GSE21927\_SPLEEN\_MONOCYTE\_VS\_GMCSF\_GCSF\_BONE\_MARROW\_UP  
GSE37605\_NOD\_VS\_C57BL6\_IRES\_GFP\_TREG\_UP, GSE37605\_NOD\_VS\_C57BL6\_IRES\_GFP\_TREG\_UP  
IWANAGA\_CARCINOGENESIS\_BY\_KRAS\_UP, IWANAGA\_CARCINOGENESIS\_BY\_KRAS\_UP  
CCAGGGG\_MIR331, CCAGGGG\_MIR331  
GSE38697\_LIGHT\_ZONE\_VS\_DARK\_ZONE\_BCELL\_DN, GSE38697\_LIGHT\_ZONE\_VS\_DARK\_ZONE\_BCELL\_DN  
GOBP\_NEGATIVE\_REGULATION\_OF\_MYOTUBE\_DIFFERENTIATION, GOBP\_NEGATIVE\_REGULATION\_OF\_MYOTUBE\_DIFFERENTIATION