


**NAIVE\_CD4\_TCELL\_CORD\_BLOOD\_UP, GSE1460\_DP\_THYMOCYTE\_VS\_NAIVE\_CD4\_TCELL\_CORD\_BLOOD\_UP**



GSE13738\_TCR\_VS\_BYSTANDER\_ACTIVATED\_CD4\_TCELL\_UP, GSE13738\_TCR\_VS\_BYSTANDER\_ACTIVATED\_CD4\_  
GSE37605\_C57BL6\_VS\_NOD\_FOXP3\_IRES\_GFP\_TCONV\_DN, GSE37605\_C57BL6\_VS\_NOD\_FOXP3\_IRES\_GFP\_TCONV\_  
GSE3920\_UNTREATED\_VS\_IFNB\_TREATED\_ENDOTHELIAL\_CELL\_UP, GSE3920\_UNTREATED\_VS\_IFNB\_TREATED\_  
REACTOME\_PI\_METABOLISM, REACTOME\_PI\_METABOLISM  
MODULE\_493, MODULE\_493  
GO\_PDZ\_DOMAIN\_BINDING, GO\_PDZ\_DOMAIN\_BINDING  
GO\_NEGATIVE\_REGULATION\_OF\_INTERLEUKIN\_2\_PRODUCTION, GO\_NEGATIVE\_REGULATION\_OF\_INTERLEU  
GO\_MISFOLDED\_PROTEIN\_BINDING, GO\_MISFOLDED\_PROTEIN\_BINDING  
chr9q22, chr9q22  
CAGGTCC\_MIR492, CAGGTCC\_MIR492