

ED\_RNA\_VS\_MRNA\_DC\_UP, GSE14000\_TRANSLATED\_RNA\_VS\_MRNA\_DC\_UP

- GSE45739\_NRAS\_KO\_VS\_WT\_ACD3\_ACD28\_STIM\_CD4\_TCELL\_UP, GSE45739\_NRAS\_KO\_VS\_WT\_ACD3\_ACD28\_STIM\_CD4\_TCELL\_UP
- GSE11864\_UNTREATED\_VS\_CSF1\_IN\_MAC\_DN, GSE11864\_UNTREATED\_VS\_CSF1\_IN\_MAC\_DN
- GSE2770\_IL12\_ACT\_VS\_ACT\_CD4\_TCELL\_2H\_DN, GSE2770\_IL12\_ACT\_VS\_ACT\_CD4\_TCELL\_2H\_DN
- GSE14000\_UNSTIM\_VS\_16H\_LPS\_DC\_DN, GSE14000\_UNSTIM\_VS\_16H\_LPS\_DC\_DN
- GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_48H\_UP, GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_48H\_UP
- GSE2770\_IL12\_AND\_TGFB\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_2H\_UP, GSE2770\_IL12\_AND\_TGFB\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_2H\_UP
- GSE19401\_PLN\_VS\_PEYERS\_PATCH\_FOLLICULAR\_DC\_DN, GSE19401\_PLN\_VS\_PEYERS\_PATCH\_FOLLICULAR\_DC\_DN
- GSE4748\_CTRL\_VS\_CYANOBACTERIUM\_LPSLIKE\_STIM\_DC\_1H\_UP, GSE4748\_CTRL\_VS\_CYANOBACTERIUM\_LPSLIKE\_STIM\_DC\_1H\_UP
- GSE1925\_CTRL\_VS\_IFNG\_PRIMED\_MACROPHAGE\_24H\_IFNG\_STIM\_DN, GSE1925\_CTRL\_VS\_IFNG\_PRIMED\_MACROPHAGE\_24H\_IFNG\_STIM\_DN
- GSE17721\_PAM3CSK4\_VS\_CPG\_0.5H\_BMDC\_UP, GSE17721\_PAM3CSK4\_VS\_CPG\_0.5H\_BMDC\_UP
- HP\_ABNORMALITY\_OF\_COAGULATION, HP\_ABNORMALITY\_OF\_COAGULATION
- MIR4277, MIR4277
- PAX5\_01, PAX5\_01
- AKT\_UP\_MTOR\_DN.V1\_DN, AKT\_UP\_MTOR\_DN.V1\_DN
- GSE23505\_UNTREATED\_VS\_4DAY\_IL6\_IL1\_TREATED\_CD4\_TCELL\_DN, GSE23505\_UNTREATED\_VS\_4DAY\_IL6\_IL1\_TREATED\_CD4\_TCELL\_DN
- MIR2117, MIR2117
- UBE2I\_TARGET\_GENES, UBE2I\_TARGET\_GENES
- MIR5579\_5P, MIR5579\_5P
- GOCC\_INO80\_TYPE\_COMPLEX, GOCC\_INO80\_TYPE\_COMPLEX
- CDX1\_TARGET\_GENES, CDX1\_TARGET\_GENES
- HP\_FRONTAL\_CORTICAL\_ATROPHY, HP\_FRONTAL\_CORTICAL\_ATROPHY
- GOBP\_POSITIVE\_REGULATION\_OF\_SMOOTHENED\_SIGNALING\_PATHWAY, GOBP\_POSITIVE\_REGULATION\_OF\_SMOOTHENED\_SIGNALING\_PATHWAY
- REACTOME\_GLYCOGEN\_SYNTHESIS, REACTOME\_GLYCOGEN\_SYNTHESIS
- WP\_EICOSANOID\_METABOLISM\_VIA\_LIPO\_OXYGENASES\_LOX, WP\_EICOSANOID\_METABOLISM\_VIA\_LIPO\_OXYGENASES\_LOX
- HP\_IMPAIRED\_TANDEM\_GAIT, HP\_IMPAIRED\_TANDEM\_GAIT
- MIR4515, MIR4515
- GSE37301\_HEMATOPOIETIC\_STEM\_CELL\_VS\_CD4\_TCELL\_DN, GSE37301\_HEMATOPOIETIC\_STEM\_CELL\_VS\_CD4\_TCELL\_DN
- YUAN\_ZNF143\_PARTNERS, YUAN\_ZNF143\_PARTNERS
- MARSON\_FOXP3\_TARGETS\_STIMULATED\_UP, MARSON\_FOXP3\_TARGETS\_STIMULATED\_UP
- WP\_GLUCURONIDATION, WP\_GLUCURONIDATION
- GOBP\_VENTRICULAR\_SYSTEM\_DEVELOPMENT, GOBP\_VENTRICULAR\_SYSTEM\_DEVELOPMENT
- GOERING\_BLOOD\_HDL\_CHOLESTEROL\_QTL\_TRANS, GOERING\_BLOOD\_HDL\_CHOLESTEROL\_QTL\_TRANS
- HP\_HYPERCHOLESTEROLEMIA, HP\_HYPERCHOLESTEROLEMIA
- MIR3914, MIR3914
- MIR3146, MIR3146
- HP\_DYSTROPHIC\_FINGERNAILS, HP\_DYSTROPHIC\_FINGERNAILS
- WP\_NONHOMOLOGOUS\_END\_JOINING, WP\_NONHOMOLOGOUS\_END\_JOINING
- GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_POSTSYNAPTIC\_MEMBRANE, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_POS
- MIR3620\_3P, MIR3620\_3P
- REACTOME\_PROSTANOID\_LIGAND\_RECEPTORS, REACTOME\_PROSTANOID\_LIGAND\_RECEPTORS
- TURASHVILI\_BREAST\_NORMAL\_DUCTAL\_VS\_LOBULAR\_DN, TURASHVILI\_BREAST\_NORMAL\_DUCTAL\_VS\_LOBULAR\_DN
- GOBP\_NEUROTRANSMITTER\_RECEPTOR\_TRANSPORT\_TO\_PLASMA\_MEMBRANE, GOBP\_NEUROTRANSMITTER\_RECEPTOR\_TRANSPORT\_TO\_PLASMA\_ME
- HP\_ABNORMAL\_SHARPEY\_FIBER\_MORPHOLOGY, HP\_ABNORMAL\_SHARPEY\_FIBER\_MORPHOLOGY