

A2AR\_AGONIST\_TREATED\_TREG\_UP, GSE34006\_UNTREATED\_VS\_A2AR\_AGONIST\_TREATED\_TREG\_UP

GSE14308\_TH1\_VS\_NAIVE\_CD4\_TCELL\_UP, GSE14308\_TH1\_VS\_NAIVE\_CD4\_TCELL\_UP  
GSE411\_UNSTIM\_VS\_100MIN\_IL6\_STIM\_MACROPHAGE\_UP, GSE411\_UNSTIM\_VS\_100MIN\_IL6\_STIM\_MACROPHAGE\_UP  
GSE23568\_ID3\_TRANSDUCED\_VS\_ID3\_KO\_CD8\_TCELL\_UP, GSE23568\_ID3\_TRANSDUCED\_VS\_ID3\_KO\_CD8\_TCELL\_UP  
GSE339\_EX\_VIVO\_VS\_IN\_CULTURE\_CD4CD8DN\_DC\_DN, GSE339\_EX\_VIVO\_VS\_IN\_CULTURE\_CD4CD8DN\_DC\_DN  
GSE26030\_TH1\_VS\_TH17\_DAY15\_POST\_POLARIZATION\_UP, GSE26030\_TH1\_VS\_TH17\_DAY15\_POST\_POLARIZATION\_UP  
GSE12198\_NK\_VS\_NK\_ACT\_EXPANSION\_SYSTEM\_DERIVED\_NK\_CELL\_UP, GSE12198\_NK\_VS\_NK\_ACT\_EXPANSION\_SYSTEM\_DERIVED\_NK\_CELL\_UP  
GSE25088\_IL4\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_MACROPHAGE\_DAY10\_UP, GSE25088\_IL4\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_MACROPHAGE\_DAY10\_UP  
EGFR\_UP.V1\_DN, EGFR\_UP.V1\_DN  
ACACTAC\_MIR1423P, ACACTAC\_MIR1423P  
GSE21360\_SECONDARY\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_DN, GSE21360\_SECONDARY\_VS\_QUATERNARY\_MEMORY\_CD8\_TCELL\_DN  
MIR1184, MIR1184  
KEGG\_FC\_GAMMA\_R\_MEDIATED\_PHAGOCYTOSIS, KEGG\_FC\_GAMMA\_R\_MEDIATED\_PHAGOCYTOSIS  
GSE21927\_UNTREATED\_VS\_GMCSF\_GCSF\_TREATED\_BONE\_MARROW\_DN, GSE21927\_UNTREATED\_VS\_GMCSF\_GCSF\_TREATED\_BONE\_MARROW\_DN  
MIR2681\_5P, MIR2681\_5P  
GOBP\_PEPTIDYL\_THREONINE\_MODIFICATION, GOBP\_PEPTIDYL\_THREONINE\_MODIFICATION  
GSE15330\_HSC\_VS\_LYMPHOID\_PRIMED\_MULTIPOTENT\_PROGENITOR\_UP, GSE15330\_HSC\_VS\_LYMPHOID\_PRIMED\_MULTIPOTENT\_PROGENITOR\_UP  
GSE2585\_THYMIC\_MACROPHAGE\_VS\_MTEC\_UP, GSE2585\_THYMIC\_MACROPHAGE\_VS\_MTEC\_UP  
GSE15735\_2H\_VS\_12H\_HDAC\_INHIBITOR\_TREATED\_CD4\_TCELL\_UP, GSE15735\_2H\_VS\_12H\_HDAC\_INHIBITOR\_TREATED\_CD4\_TCELL\_UP  
GSE36527\_CD62L\_HIGH\_CD69\_NEG\_VS\_CD62L\_LOW\_CD69\_POS\_TREG\_KLRG1\_NEG\_DN, GSE36527\_CD62L\_HIGH\_CD69\_NEG\_VS\_CD62L\_LOW\_CD69\_POS\_TREG\_KLRG1\_NEG\_DN  
MIR4704\_3P, MIR4704\_3P  
MIR9500, MIR9500  
MIR4277, MIR4277  
GOBP\_MEMBRANE\_FUSION, GOBP\_MEMBRANE\_FUSION  
IWANAGA\_CARCIANOGENESIS\_BY\_KRAS\_PTEN\_UP, IWANAGA\_CARCIANOGENESIS\_BY\_KRAS\_PTEN\_UP  
MIR1279, MIR1279  
GSE17721\_CPG\_VS\_GARDIQUIMOD\_2H\_BMDC\_UP, GSE17721\_CPG\_VS\_GARDIQUIMOD\_2H\_BMDC\_UP  
CDH4\_TARGET\_GENES, CDH4\_TARGET\_GENES  
GSE23505\_UNTREATED\_VS\_4DAY\_IL6\_IL1\_TREATED\_CD4\_TCELL\_DN, GSE23505\_UNTREATED\_VS\_4DAY\_IL6\_IL1\_TREATED\_CD4\_TCELL\_DN  
GOBP\_MODULATION\_OF\_PROCESS\_OF\_OTHER\_ORGANISM\_INVOLVED\_IN\_SYMBIOTIC\_INTERACTION, GOBP\_MODULATION\_OF\_PROCESS\_OF\_OTHER\_ORGANISM\_INVOLVED\_IN\_SYMBIOTIC\_INTERACTION  
GSE2585\_THYMIC\_DC\_VS\_THYMIC\_MACROPHAGE\_UP, GSE2585\_THYMIC\_DC\_VS\_THYMIC\_MACROPHAGE\_UP  
GSE17721\_CPG\_VS\_GARDIQUIMOD\_12H\_BMDC\_UP, GSE17721\_CPG\_VS\_GARDIQUIMOD\_12H\_BMDC\_UP  
MIR4314, MIR4314  
MIR6882\_3P, MIR6882\_3P  
GOBP\_MODULATION\_OF\_PROCESS\_OF\_OTHER\_ORGANISM, GOBP\_MODULATION\_OF\_PROCESS\_OF\_OTHER\_ORGANISM  
MIR593\_3P, MIR593\_3P  
WP\_INTEGRINMEDIATED\_CELL\_ADHESION, WP\_INTEGRINMEDIATED\_CELL\_ADHESION  
GSE15750\_WT\_VS\_TRAF6KO\_DAY6\_EFF\_CD8\_TCELL\_DN, GSE15750\_WT\_VS\_TRAF6KO\_DAY6\_EFF\_CD8\_TCELL\_DN  
DORSAM\_HOXA9\_TARGETS\_UP, DORSAM\_HOXA9\_TARGETS\_UP  
HP\_WADDLING\_GAIT, HP\_WADDLING\_GAIT  
GOBP\_REGULATION\_OF\_AUTOPHAGY\_OF\_MITOCHONDRION, GOBP\_REGULATION\_OF\_AUTOPHAGY\_OF\_MITOCHONDRION  
GOBP\_EXOCYTIC\_PROCESS, GOBP\_EXOCYTIC\_PROCESS  
GOCC\_PROTEASOME\_REGULATORY\_PARTICLE\_LID\_SUBCOMPLEX, GOCC\_PROTEASOME\_REGULATORY\_PARTICLE\_LID\_SUBCOMPLEX  
MIR3654, MIR3654  
GOBP\_VESICLE\_DOCKING\_INVOLVED\_IN\_EXOCYTOSIS, GOBP\_VESICLE\_DOCKING\_INVOLVED\_IN\_EXOCYTOSIS  
LIANG\_HEMATOPOIESIS\_STEM\_CELL\_NUMBER\_SMALL\_VS\_HUGE\_UP, LIANG\_HEMATOPOIESIS\_STEM\_CELL\_NUMBER\_SMALL\_VS\_HUGE\_UP  
CDPCR1\_01, CDPCR1\_01  
MIR193B\_5P, MIR193B\_5P  
MIR6515\_5P, MIR6515\_5P  
GOBP\_GLANDULAR\_EPITHELIAL\_CELL\_DIFFERENTIATION, GOBP\_GLANDULAR\_EPITHELIAL\_CELL\_DIFFERENTIATION  
GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_THE\_CH\_NH\_GROUP\_OF\_DONORS\_NAD\_OR\_NADP\_AS\_ACCEPTOR, GOMF\_OXIDOREDUCTASE\_ACTIVITY\_ACTING\_ON\_THE\_CH\_NH\_GROUP\_OF\_DONORS\_NAD\_OR\_NADP\_AS\_ACCEPTOR  
GOBP\_COLUMNAR\_CUBOIDAL\_EPITHELIAL\_CELL\_DIFFERENTIATION, GOBP\_COLUMNAR\_CUBOIDAL\_EPITHELIAL\_CELL\_DIFFERENTIATION  
YAATNANRNNNCAG\_UNKNOWN, YAATNANRNNNCAG\_UNKNOWN  
GOBP\_ACTIN\_FILAMENT\_SEVERING, GOBP\_ACTIN\_FILAMENT\_SEVERING  
chr16q13, chr16q13