

ND\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP, GSE5542\_IFNG\_VS\_IFNA\_AND\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP

GSE27786\_ERYTHROBLAST\_VS\_NEUTROPHIL\_DN, GSE27786\_ERYTHROBLAST\_VS\_NEUTROPHIL\_DN  
GSE5542\_UNTREATED\_VS\_IFNA\_AND\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP, GSE5542\_UNTREATED\_VS\_IFNA\_AND\_IFNG\_TREATED\_EPITHELIAL\_CELLS\_24H\_UP  
GSE27786\_LIN\_NEG\_VS\_NKCELL\_DN, GSE27786\_LIN\_NEG\_VS\_NKCELL\_DN  
GSE11961\_MEMORY\_BCELL\_DAY7\_VS\_GERMINAL\_CENTER\_BCELL\_DAY40\_DN, GSE11961\_MEMORY\_BCELL\_DAY7\_VS\_GERMINAL\_CENTER\_BCELL\_DAY40\_DN  
GSE45881\_CXCR6HI\_VS\_CXCR1LO\_COLONIC\_LAMINA\_PROPRIA\_DN, GSE45881\_CXCR6HI\_VS\_CXCR1LO\_COLONIC\_LAMINA\_PROPRIA\_DN  
GSE11961\_GERMINAL\_CENTER\_BCELL\_DAY7\_VS\_MEMORY\_BCELL\_DAY40\_UP, GSE11961\_GERMINAL\_CENTER\_BCELL\_DAY7\_VS\_MEMORY\_BCELL\_DAY40\_UP  
GSE27786\_LSK\_VS\_MONO\_MAC\_DN, GSE27786\_LSK\_VS\_MONO\_MAC\_DN  
GSE28737\_FOLLICULAR\_VS\_MARGINAL\_ZONE\_BCELL\_DN, GSE28737\_FOLLICULAR\_VS\_MARGINAL\_ZONE\_BCELL\_DN  
GSE44732\_UNSTIM\_VS\_IL27\_STIM\_IMATURE\_DC\_UP, GSE44732\_UNSTIM\_VS\_IL27\_STIM\_IMATURE\_DC\_UP  
GSE14308\_TH17\_VS\_NAIVE\_CD4\_TCELL\_DN, GSE14308\_TH17\_VS\_NAIVE\_CD4\_TCELL\_DN  
MIR8057, MIR8057  
GSE18893\_CTRL\_VS\_TNF\_TREATED\_TREG\_24H\_UP, GSE18893\_CTRL\_VS\_TNF\_TREATED\_TREG\_24H\_UP  
GSE5542\_IFNG\_VS\_IFNA\_TREATED\_EPITHELIAL\_CELLS\_6H\_DN, GSE5542\_IFNG\_VS\_IFNA\_TREATED\_EPITHELIAL\_CELLS\_6H\_DN  
CCAWNWWNNNGGC\_UNKNOWN, CCAWNWWNNNGGC\_UNKNOWN  
GSE17186\_NAIVE\_VS\_CD21HIGH\_TRANSITIONAL\_BCELL\_UP, GSE17186\_NAIVE\_VS\_CD21HIGH\_TRANSITIONAL\_BCELL\_UP  
MODULE\_567, MODULE\_567  
GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_GREY\_DN, GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_GREY\_DN  
MIR6818\_5P, MIR6818\_5P  
WEIGEL\_OXIDATIVE\_STRESS\_BY\_HNE\_AND\_TBH, WEIGEL\_OXIDATIVE\_STRESS\_BY\_HNE\_AND\_TBH  
HP\_LIPODYSTROPHY, HP\_LIPODYSTROPHY  
GOMF\_MYOSIN\_BINDING, GOMF\_MYOSIN\_BINDING  
GOBP\_VIRAL\_BUDDING, GOBP\_VIRAL\_BUDDING  
GOBP\_POLYSACCHARIDE\_METABOLIC\_PROCESS, GOBP\_POLYSACCHARIDE\_METABOLIC\_PROCESS  
GOBP\_POSITIVE\_REGULATION\_OF\_DOUBLE\_STRAND\_BREAK\_REPAIR, GOBP\_POSITIVE\_REGULATION\_OF\_DOUBLE\_STRAND\_BREAK\_REPAIR  
ZNF169\_TARGET\_GENES, ZNF169\_TARGET\_GENES  
HP\_ABNORMAL\_BRONCHUS\_MORPHOLOGY, HP\_ABNORMAL\_BRONCHUS\_MORPHOLOGY  
MIR6888\_3P, MIR6888\_3P  
MIR12114, MIR12114  
MARIADASON\_REGULATED\_BY\_HISTONE\_ACETYLATION\_DN, MARIADASON\_REGULATED\_BY\_HISTONE\_ACETYLATION\_DN  
PID\_ECADHERIN\_KERATINOCYTE\_PATHWAY, PID\_ECADHERIN\_KERATINOCYTE\_PATHWAY  
MIR6864\_5P, MIR6864\_5P  
MOSERLE\_IFNA\_RESPONSE, MOSERLE\_IFNA\_RESPONSE  
REACTOME\_FGFR1\_MUTANT\_RECEPTOR\_ACTIVATION, REACTOME\_FGFR1\_MUTANT\_RECEPTOR\_ACTIVATION  
HP\_ABNORMAL\_MUSCLE\_GLYCOGEN\_CONTENT, HP\_ABNORMAL\_MUSCLE\_GLYCOGEN\_CONTENT  
HP\_CARCINOMA, HP\_CARCINOMA  
BOWIE\_RESPONSE\_TO\_TAMOXIFEN, BOWIE\_RESPONSE\_TO\_TAMOXIFEN  
GOBP\_CELLULAR\_GLUCAN\_METABOLIC\_PROCESS, GOBP\_CELLULAR\_GLUCAN\_METABOLIC\_PROCESS  
HP\_HIGH\_MYOPIA, HP\_HIGH\_MYOPIA  
REACTOME\_GLYCOGEN\_BREAKDOWN\_GLYCOGENOLYSIS, REACTOME\_GLYCOGEN\_BREAKDOWN\_GLYCOGENOLYSIS  
GOBP\_XENOBIOTIC\_TRANSPORT, GOBP\_XENOBIOTIC\_TRANSPORT  
DORN\_ADENOVIRUS\_INFECTION\_12HR\_UP, DORN\_ADENOVIRUS\_INFECTION\_12HR\_UP  
BOWIE\_RESPONSE\_TO\_EXTRACELLULAR\_MATRIX, BOWIE\_RESPONSE\_TO\_EXTRACELLULAR\_MATRIX  
MIR3690, MIR3690  
GU\_PDEF\_TARGETS\_UP, GU\_PDEF\_TARGETS\_UP  
ZHANG\_RESPONSE\_TO\_CANTHARIDIN\_UP, ZHANG\_RESPONSE\_TO\_CANTHARIDIN\_UP  
GNF2\_IGFBP1, GNF2\_IGFBP1  
MIR6802\_5P, MIR6802\_5P  
BIOCARTA\_IFNA\_PATHWAY, BIOCARTA\_IFNA\_PATHWAY  
TSAI\_DNAJB4\_TARGETS\_UP, TSAI\_DNAJB4\_TARGETS\_UP  
GNF2\_TIMP2, GNF2\_TIMP2  
REACTOME\_SYNTHESIS\_OF\_PYROPHOSPHATES\_IN\_THE\_CYTOSOL, REACTOME\_SYNTHESIS\_OF\_PYROPHOSPHATES\_IN\_THE\_CYTOSOL  
GNF2\_KISS1, GNF2\_KISS1  
MIR4695\_3P, MIR4695\_3P  
XU\_RESPONSE\_TO\_TRETINOIN\_AND\_NSC682994\_UP, XU\_RESPONSE\_TO\_TRETINOIN\_AND\_NSC682994\_UP  
HP\_REDUCED\_FORCED\_EXPIRATORY\_VOLUME\_IN\_ONE\_SECOND, HP\_REDUCED\_FORCED\_EXPIRATORY\_VOLUME\_IN\_ONE\_SECOND  
WP\_IL10\_ANTIINFLAMMATORY\_SIGNALING\_PATHWAY, WP\_IL10\_ANTIINFLAMMATORY\_SIGNALING\_PATHWAY  
HP\_PEDAL\_EDEMA, HP\_PEDAL\_EDEMA  
MIR3678\_5P, MIR3678\_5P  
GOBP\_LEFT\_RIGHT\_PATTERN\_FORMATION, GOBP\_LEFT\_RIGHT\_PATTERN\_FORMATION