

4\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_TGFB\_AND\_IL4\_VS\_IL4\_TREATED\_ACT\_CD4\_TCELL\_6H\_UP

GSE2770\_TGFB\_AND\_IL4\_IL12\_TREATED\_ACT\_CD4\_TCELL\_48H\_UP, GSE2770\_TGFB\_AND\_IL4\_IL12\_TREATED\_ACT\_CD4\_TCELL\_48H\_UP  
GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_6H\_UP, GSE2770\_TGFB\_AND\_IL4\_ACT\_VS\_ACT\_CD4\_TCELL\_6H\_UP  
MODULE\_257, MODULE\_257  
GSE22886\_NAIVE\_CD4\_TCELL\_VS\_NKCELL\_UP, GSE22886\_NAIVE\_CD4\_TCELL\_VS\_NKCELL\_UP  
HP\_SINGLE\_TRANSVERSE\_PALMAR\_CREASE, HP\_SINGLE\_TRANSVERSE\_PALMAR\_CREASE  
GSE45365\_HEALTHY\_VS\_MCMV\_INFECTION\_CD11B\_DC\_IFNAR\_KO\_UP, GSE45365\_HEALTHY\_VS\_MCMV\_INFECTION\_CD11B\_DC\_IFNAR\_KO\_UP  
MIR4682, MIR4682  
HP\_NYCTALOPIA, HP\_NYCTALOPIA  
HP\_ECZEMA, HP\_ECZEMA  
HP\_FLAT\_FACE, HP\_FLAT\_FACE  
GOBP\_STRESS\_FIBER\_ASSEMBLY, GOBP\_STRESS\_FIBER\_ASSEMBLY  
GSE16385\_IFNG\_TNF\_VS\_IL4\_STIM\_MACROPHAGE\_ROSIGLITAZONE\_TREATED\_UP, GSE16385\_IFNG\_TNF\_VS\_IL4\_STIM\_MACROPHAGE\_ROSIGLITAZONE\_TREATED\_UP  
GOBP\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY, GOBP\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY  
HP\_ABNORMALITY\_OF\_THE\_INCISOR, HP\_ABNORMALITY\_OF\_THE\_INCISOR  
GOBP\_REGULATION\_OF\_ACTOMYOSIN\_STRUCTURE\_ORGANIZATION, GOBP\_REGULATION\_OF\_ACTOMYOSIN\_STRUCTURE\_ORGANIZATION  
HP\_PROGRESSIVE\_NIGHT\_BLINDNESS, HP\_PROGRESSIVE\_NIGHT\_BLINDNESS  
WP\_PATHWAYS\_REGULATING\_HIPPO\_SIGNALING, WP\_PATHWAYS\_REGULATING\_HIPPO\_SIGNALING  
HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_RIBS, HP\_APLASIA\_HYPOPLASIA\_OF\_THE\_RIBS  
GOBP\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY, GOBP\_POSITIVE\_REGULATION\_OF\_ACTIN\_FILAMENT\_BUNDLE\_ASSEMBLY  
MIR6895\_3P, MIR6895\_3P  
GOBP\_REGULATION\_OF\_SIGNALING\_RECEPTOR\_ACTIVITY, GOBP\_REGULATION\_OF\_SIGNALING\_RECEPTOR\_ACTIVITY  
GOMF\_RNA\_POLYMERASE\_CORE\_ENZYME\_BINDING, GOMF\_RNA\_POLYMERASE\_CORE\_ENZYME\_BINDING  
GOMF\_ION\_CHANNEL\_REGULATOR\_ACTIVITY, GOMF\_ION\_CHANNEL\_REGULATOR\_ACTIVITY  
HP\_RETINAL\_ATROPHY, HP\_RETINAL\_ATROPHY  
YU\_MYC\_TARGETS\_DN, YU\_MYC\_TARGETS\_DN  
SRPK2\_TARGET\_GENES, SRPK2\_TARGET\_GENES  
GOBP\_REGULATION\_OF\_HISTONE\_H4\_ACETYLTATION, GOBP\_REGULATION\_OF\_HISTONE\_H4\_ACETYLTATION  
LIAN\_NEUTROPHIL\_GANULE\_CONSTITUENTS, LIAN\_NEUTROPHIL\_GANULE\_CONSTITUENTS  
HP\_HYPOPLASTIC\_FACIAL\_BONES, HP\_HYPOPLASTIC\_FACIAL\_BONES  
GOMF\_ENOYL\_COA\_HYDRATASE\_ACTIVITY, GOMF\_ENOYL\_COA\_HYDRATASE\_ACTIVITY  
VANHARANTA\_UTERINE\_FIBROID\_UP, VANHARANTA\_UTERINE\_FIBROID\_UP  
HP\_HEPATIC\_CYSTS, HP\_HEPATIC\_CYSTS  
DACOSTA\_ERCC3\_ALLELE\_XPCS\_VS\_TTD\_UP, DACOSTA\_ERCC3\_ALLELE\_XPCS\_VS\_TTD\_UP  
BREDEMEYER\_RAG\_SIGNALING\_VIA\_ATM\_NOT\_VIA\_NFKB\_DN, BREDEMEYER\_RAG\_SIGNALING\_VIA\_ATM\_NOT\_VIA\_NFKB\_DN  
HP\_TRANSIENT\_GLOBAL\_AMNESIA, HP\_TRANSIENT\_GLOBAL\_AMNESIA  
HP\_ABNORMAL\_NUMBER\_OF\_PERMANENT\_TEETH, HP\_ABNORMAL\_NUMBER\_OF\_PERMANENT\_TEETH  
MIR6810\_5P, MIR6810\_5P  
HP\_RENAL\_FIBROSIS, HP\_RENAL\_FIBROSIS  
GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_POSTSYNAPTIC\_MEMBRANE, GOBP\_ESTABLISHMENT\_OF\_PROTEIN\_LOCALIZATION\_TO\_POSTSYNAPTIC\_MEMBRANE  
HP\_CEREBELLAR\_DYSPLASIA, HP\_CEREBELLAR\_DYSPLASIA  
GOBP\_LATE\_ENDOSOME\_TO\_VACUOLE\_TRANSPORT\_VIA\_MULTIVESICULAR\_BODY\_SORTING\_PATHWAY, GOBP\_LATE\_ENDOSOME\_TO\_VACUOLE\_TRANSPORT\_VIA\_MULTIVESICULAR\_BODY\_SORTING\_PATHWAY  
HP\_NARROW\_NOSE, HP\_NARROW\_NOSE  
HP\_ATELECTASIS, HP\_ATELECTASIS  
GOBP\_NEUROTRANSMITTER\_RECEPTOR\_TRANSPORT\_TO\_PLASMA\_MEMBRANE, GOBP\_NEUROTRANSMITTER\_RECEPTOR\_TRANSPORT\_TO\_PLASMA\_MEMBRANE  
GOMF\_BETA\_1\_3\_GALACTOSYLTRANSFERASE\_ACTIVITY, GOMF\_BETA\_1\_3\_GALACTOSYLTRANSFERASE\_ACTIVITY  
GOBP\_CYSTEINE\_METABOLIC\_PROCESS, GOBP\_CYSTEINE\_METABOLIC\_PROCESS  
GOBP\_PROTEIN\_LOCALIZATION\_TO\_SYNAPSE, GOBP\_PROTEIN\_LOCALIZATION\_TO\_SYNAPSE  
REACTOME\_UPTAKE\_AND\_FUNCTION\_OF\_ANTHRAX\_TOXINS, REACTOME\_UPTAKE\_AND\_FUNCTION\_OF\_ANTHRAX\_TOXINS  
GOBP\_REGULATION\_OF\_FATTY\_ACID\_BETA\_OXIDATION, GOBP\_REGULATION\_OF\_FATTY\_ACID\_BETA\_OXIDATION  
MODULE\_431, MODULE\_431  
HP\_HYPOPIGMENTATION\_OF\_THE\_FUNDUS, HP\_HYPOPIGMENTATION\_OF\_THE\_FUNDUS  
WP\_1Q211\_COPY\_NUMBER\_VARIATION\_SYNDROME, WP\_1Q211\_COPY\_NUMBER\_VARIATION\_SYNDROME  
GOBP\_NEGATIVE\_REGULATION\_OF\_ALPHA\_BETA\_T\_CELL\_PROLIFERATION, GOBP\_NEGATIVE\_REGULATION\_OF\_ALPHA\_BETA\_T\_CELL\_PROLIFERATION  
HP\_Y\_SHAPED\_METACARPALS, HP\_Y\_SHAPED\_METACARPALS  
HP\_ABNORMALITY\_OF\_FUNDUS\_PIGMENTATION, HP\_ABNORMALITY\_OF\_FUNDUS\_PIGMENTATION  
GOCC\_POSTSYNAPTIC\_ENDOSOME, GOCC\_POSTSYNAPTIC\_ENDOSOME  
GOCC\_POSTSYNAPTIC\_DENSITY\_MEMBRANE, GOCC\_POSTSYNAPTIC\_DENSITY\_MEMBRANE  
GOBP\_ENDOSOME\_TO\_PLASMA\_MEMBRANE\_PROTEIN\_TRANSPORT, GOBP\_ENDOSOME\_TO\_PLASMA\_MEMBRANE\_PROTEIN\_TRANSPORT  
GOBP\_EMBRYONIC\_CAMERA\_TYPE\_EYE\_FORMATION, GOBP\_EMBRYONIC\_CAMERA\_TYPE\_EYE\_FORMATION  
chr3q13, chr3q13  
GOBP\_XENOBIOTIC\_CATABOLIC\_PROCESS, GOBP\_XENOBIOTIC\_CATABOLIC\_PROCESS