

IAL\_GENE\_EXPRESSION, GOBP\_MITOCHONDRIAL\_GENE\_EXPRESSION

- GOBP\_TRNA\_METABOLIC\_PROCESS, GOBP\_TRNA\_METABOLIC\_PROCESS
- RAMALHO\_STEMNESS\_UP, RAMALHO\_STEMNESS\_UP
- GSE17322\_CD103\_POS\_VS\_CD11B\_HIGH\_LUNG\_DC\_DN, GSE17322\_CD103\_POS\_VS\_CD11B\_HIGH\_LUNG\_DC\_DN
- HP\_ABNORMALITY\_OF\_THE\_MITOCHONDRION, HP\_ABNORMALITY\_OF\_THE\_MITOCHONDRION
- GOMF\_RIBONUCLEOPROTEIN\_COMPLEX\_BINDING, GOMF\_RIBONUCLEOPROTEIN\_COMPLEX\_BINDING
- GSE15930\_STIM\_VS\_STIM\_AND\_IFNAB\_72H\_CD8\_T\_CELL\_DN, GSE15930\_STIM\_VS\_STIM\_AND\_IFNAB\_72H\_CD8\_T\_CELL\_DN
- GSE25123\_ROSIGLITAZONE\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_PPARG\_KO\_MACROPHAGE\_DAY10\_UP, GSE25123\_ROSIGLITAZONE\_VS\_IL4\_AND\_ROSIGLITAZONE\_STIM\_PPARG\_KO\_MACROPHAGE\_DAY10\_UP
- GOBP\_TRNA\_PROCESSING, GOBP\_TRNA\_PROCESSING
- GOMF\_CATALYTIC\_ACTIVITY\_ACTING\_ON\_A\_TRNA, GOMF\_CATALYTIC\_ACTIVITY\_ACTING\_ON\_A\_TRNA
- GOBP\_RNA\_MODIFICATION, GOBP\_RNA\_MODIFICATION
- GOMF\_LIGASE\_ACTIVITY, GOMF\_LIGASE\_ACTIVITY
- GSE20198\_UNTREATED\_VS\_IL12\_IL18\_TREATED\_ACT\_CD4\_TCELL\_DN, GSE20198\_UNTREATED\_VS\_IL12\_IL18\_TREATED\_ACT\_CD4\_TCELL\_DN
- GSE13306\_TREG\_VS\_TCONV\_DN, GSE13306\_TREG\_VS\_TCONV\_DN
- GSE20484\_MCSG\_VS\_CXCL4\_MONOCYTE\_DERIVED\_MACROPHAGE\_UP, GSE20484\_MCSG\_VS\_CXCL4\_MONOCYTE\_DERIVED\_MACROPHAGE\_UP
- GOBP\_MITOCHONDRIAL\_RESPIRATORY\_CHAIN\_COMPLEX\_ASSEMBLY, GOBP\_MITOCHONDRIAL\_RESPIRATORY\_CHAIN\_COMPLEX\_ASSEMBLY
- GSE27786\_CD8\_TCELL\_VS\_NKCELL\_UP, GSE27786\_CD8\_TCELL\_VS\_NKCELL\_UP
- GSE32423\_IL7\_VS\_IL4\_MEMORY\_CD8\_TCELL\_DN, GSE32423\_IL7\_VS\_IL4\_MEMORY\_CD8\_TCELL\_DN
- GSE25087\_TREG\_VS\_TCONV\_ADULT\_DN, GSE25087\_TREG\_VS\_TCONV\_ADULT\_DN
- HP\_DELAYED\_MYELINATION, HP\_DELAYED\_MYELINATION
- MOREAUX\_B\_LYMPHOCYTE\_MATURATION\_BY\_TACI\_DN, MOREAUX\_B\_LYMPHOCYTE\_MATURATION\_BY\_TACI\_DN
- HP\_ABNORMAL\_ACTIVITY\_OF\_MITOCHONDRIAL\_RESPIRATORY\_CHAIN, HP\_ABNORMAL\_ACTIVITY\_OF\_MITOCHONDRIAL\_RESPIRATORY\_CHAIN
- GSE21033\_1H\_VS\_24H\_POLYIC\_STIM\_DC\_DN, GSE21033\_1H\_VS\_24H\_POLYIC\_STIM\_DC\_DN
- GOBP\_RNA\_METHYLATION, GOBP\_RNA\_METHYLATION
- ZNF667\_TARGET\_GENES, ZNF667\_TARGET\_GENES
- GOBP\_TRNA\_MODIFICATION, GOBP\_TRNA\_MODIFICATION
- GSE17721\_LPS\_VS\_POLYIC\_0.5H\_BMDC\_UP, GSE17721\_LPS\_VS\_POLYIC\_0.5H\_BMDC\_UP
- GSE17721\_CTRL\_VS\_GARDIQUIMOD\_2H\_BMDC\_UP, GSE17721\_CTRL\_VS\_GARDIQUIMOD\_2H\_BMDC\_UP
- GSE15930\_STIM\_VS\_STIM\_AND\_TRICHOSTATINA\_72H\_CD8\_T\_CELL\_DN, GSE15930\_STIM\_VS\_STIM\_AND\_TRICHOSTATINA\_72H\_CD8\_T\_CELL\_DN
- GSE27786\_CD8\_TCELL\_VS\_NKTCELL\_UP, GSE27786\_CD8\_TCELL\_VS\_NKTCELL\_UP
- GSE5589\_WT\_VS\_IL10\_KO\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_180MIN\_UP, GSE5589\_WT\_VS\_IL10\_KO\_LPS\_AND\_IL10\_STIM\_MACROPHAGE\_180MIN\_UP
- GOMF\_TRNA\_BINDING, GOMF\_TRNA\_BINDING
- BYSTRYKH\_HEMATOPOIESIS\_STEM\_CELL\_AND\_BRAIN\_QTL\_CIS, BYSTRYKH\_HEMATOPOIESIS\_STEM\_CELL\_AND\_BRAIN\_QTL\_CIS
- GOBP\_AMINO\_ACID\_ACTIVATION, GOBP\_AMINO\_ACID\_ACTIVATION
- HP\_EXERCISE\_INTOLERANCE, HP\_EXERCISE\_INTOLERANCE
- GOBP\_MITOCHONDRIAL\_RNA\_METABOLIC\_PROCESS, GOBP\_MITOCHONDRIAL\_RNA\_METABOLIC\_PROCESS
- HP\_DISTAL\_SENSORY\_IMPAIRMENT, HP\_DISTAL\_SENSORY\_IMPAIRMENT
- GSE13522\_WT\_VS\_IFNAR\_KO\_SKING\_T\_CRUZL\_Y\_STRAIN\_INF\_DN, GSE13522\_WT\_VS\_IFNAR\_KO\_SKING\_T\_CRUZL\_Y\_STRAIN\_INF\_DN
- GOMF\_LIGASE\_ACTIVITY\_FORMING\_CARBON\_OXYGEN\_BONDS, GOMF\_LIGASE\_ACTIVITY\_FORMING\_CARBON\_OXYGEN\_BONDS
- KEGG\_AMINOACYL\_TRNA\_BIOSYNTHESIS, KEGG\_AMINOACYL\_TRNA\_BIOSYNTHESIS
- GOBP\_RRNA\_MODIFICATION, GOBP\_RRNA\_MODIFICATION
- REACTOME\_TRNA\_MODIFICATION\_IN\_THE\_NUCLEUS\_AND\_CYTOSOL, REACTOME\_TRNA\_MODIFICATION\_IN\_THE\_NUCLEUS\_AND\_CYTOSOL
- GOBP\_TRNA\_METHYLATION, GOBP\_TRNA\_METHYLATION
- HP\_IMPAIRED\_VIBRATORY\_SENSATION, HP\_IMPAIRED\_VIBRATORY\_SENSATION
- GOMF\_TRNA\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_TRNA\_METHYLTRANSFERASE\_ACTIVITY
- HP\_PROGRESSIVE\_INABILITY\_TO\_WALK, HP\_PROGRESSIVE\_INABILITY\_TO\_WALK
- GOBP\_MRNA\_MODIFICATION, GOBP\_MRNA\_MODIFICATION
- HP\_RAGGED\_RED\_MUSCLE\_FIBERS, HP\_RAGGED\_RED\_MUSCLE\_FIBERS
- HP\_PROGRESSIVE\_MUSCLE\_WEAKNESS, HP\_PROGRESSIVE\_MUSCLE\_WEAKNESS
- GOMF\_ENDORIBONUCLEASE\_ACTIVITY\_PRODUCING\_5\_PHOSPHOMONOESTERS, GOMF\_ENDORIBONUCLEASE\_ACTIVITY\_PRODUCING\_5\_PHOSPHOMONOESTERS
- GOBP\_MRNA\_METHYLATION, GOBP\_MRNA\_METHYLATION
- MIR4777\_3P, MIR4777\_3P
- REACTOME\_MITOCHONDRIAL\_TRNA\_AMINOACYLATION, REACTOME\_MITOCHONDRIAL\_TRNA\_AMINOACYLATION
- GOMF\_CARBOXYLIC\_ESTER\_HYDROLASE\_ACTIVITY, GOMF\_CARBOXYLIC\_ESTER\_HYDROLASE\_ACTIVITY
- FRANCO\_BLOOD\_SANOFI\_PASTEUR\_SA\_INACTIVATED\_INFLUENZA\_VACCINE\_CORRELATED\_WITH\_ANTIBODY\_RESPONSE\_AGE\_18\_40YO\_14DY\_POSITIVE, FRANCO\_BLOOD\_SANOFI\_PASTEUR\_SA\_INACTIVATED\_INFLUENZA\_VACCINE\_CORRELATED\_WITH\_ANTIBODY\_RESPONSE\_AG
- HP\_NEONATAL\_DEATH, HP\_NEONATAL\_DEATH
- HP\_LOSS\_OF\_ABILITY\_TO\_WALK, HP\_LOSS\_OF\_ABILITY\_TO\_WALK
- DESCARTES\_MAIN\_FETAL\_PARIETAL\_AND\_CHIEF\_CELLS, DESCARTES\_MAIN\_FETAL\_PARIETAL\_AND\_CHIEF\_CELLS
- GOBP\_RESPIRATORY\_SYSTEM\_PROCESS, GOBP\_RESPIRATORY\_SYSTEM\_PROCESS
- HP\_DISTAL\_UPPER\_LIMB\_AMYOTROPHY, HP\_DISTAL\_UPPER\_LIMB\_AMYOTROPHY
- HP\_HAMMERTOES, HP\_HAMMERTOES
- HP\_DISTAL\_LOWER\_LIMB\_MUSCLE\_WEAKNESS, HP\_DISTAL\_LOWER\_LIMB\_MUSCLE\_WEAKNESS
- GOBP\_AMINOACYL\_TRNA\_METABOLISM\_INVOLVED\_IN\_TRANSLATIONAL\_FIDELITY, GOBP\_AMINOACYL\_TRNA\_METABOLISM\_INVOLVED\_IN\_TRANSLATIONAL\_FIDELITY
- GOMF\_AMINOACYL\_TRNA\_EDITING\_ACTIVITY, GOMF\_AMINOACYL\_TRNA\_EDITING\_ACTIVITY
- GOMF\_TRNA\_GUANINE\_METHYLTRANSFERASE\_ACTIVITY, GOMF\_TRNA\_GUANINE\_METHYLTRANSFERASE\_ACTIVITY
- REACTOME\_RRNA\_PROCESSING\_IN\_THE\_MITOCHONDRION, REACTOME\_RRNA\_PROCESSING\_IN\_THE\_MITOCHONDRION
- HP\_EPILEPSIA\_PARTIALIS\_CONTINUA, HP\_EPILEPSIA\_PARTIALIS\_CONTINUA
- GOBP\_TRNA\_AMINOACYLATION\_FOR\_MITOCHONDRIAL\_PROTEIN\_TRANSLATION, GOBP\_TRNA\_AMINOACYLATION\_FOR\_MITOCHONDRIAL\_PROTEIN\_TRANSLATION
- GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_BLACK\_DN, GARGALOVIC\_RESPONSE\_TO\_OXIDIZED\_PHOSPHOLIPIDS\_BLACK\_DN
- GOBP\_REGULATION\_OF\_DENDRITIC\_CELL\_CHEMOTAXIS, GOBP\_REGULATION\_OF\_DENDRITIC\_CELL\_CHEMOTAXIS
- GOBP\_MITOCHONDRIAL\_DNA\_REPLICATION, GOBP\_MITOCHONDRIAL\_DNA\_REPLICATION
- GOBP\_MOTOR\_BEHAVIOR, GOBP\_MOTOR\_BEHAVIOR
- MIR1277\_3P, MIR1277\_3P
- HP\_FIRST\_DORSAL\_INTEROSSEI\_MUSCLE\_WEAKNESS, HP\_FIRST\_DORSAL\_INTEROSSEI\_MUSCLE\_WEAKNESS
- GOBP\_REGULATION\_OF\_RESPIRATORY\_GASEOUS\_EXCHANGE, GOBP\_REGULATION\_OF\_RESPIRATORY\_GASEOUS\_EXCHANGE