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|-----|------------|---|-----------|-----------|----|--|
| 206 | GO:008090 | regulation of primary metabolic process | 4.124E-01 | 2.175E+01 | 18 | ADAM17 CXCL10 FOXPA4 HCK IL10 IL12B MTOR NFAT5 NFkB1 NFKB1 IRG3S3 ROBO1 RPTOR STAT1 STAT6 TNSL WNT1 ZC3H12D |
| 214 | GO:0031341 | regulation of cell killing | 4.167E-01 | 2.187E+01 | 3 | CADM1 JL10 IL12B |
| 211 | GO:0051270 | regulation of cellular component movement | 4.267E-01 | 2.228E+01 | 7 | ADAM10 ADAM17 CXCL10 IL8 MTOR PLXNB1 ROBO1 |
| 212 | GO:0140131 | positive regulation of lymphocyte chemotaxis | 4.286E-01 | 2.228E+01 | 2 | ADAM10 ADAM17 |
| 214 | GO:0045088 | regulation of innate immune response | 4.378E-01 | 2.265E+01 | 5 | CADM1 HCK IL12B NFkB1 STAT1 |
| 214 | GO:0008593 | regulation of Notch signaling pathway | 4.416E-01 | 2.274E+01 | 3 | ADAM10 ROBO1 WNT1 |
| 216 | GO:0019222 | regulation of metabolic process | 4.557E-01 | 2.336E+01 | 19 | ADAM17 CXCL10 FOXPA4 HCK IL10 IL12B MTOR NFAT5 NFkB1 NFKB1 IRG3S3 ROBO1 RPTOR SLC6A4 STAT1 STAT6 TNSL WNT1 ZC3H12D |
| 216 | GO:0002708 | positive regulation of lymphocyte mediated immunity | 4.674E-01 | 2.385E+01 | 3 | CADM1 JL12B STAT6 |
| 211 | GO:0035556 | intracellular signal transduction | 4.754E-01 | 2.414E+01 | 9 | CXCL10 IL5RA IL8 MTOR NFAT5 NFkB1 PLXNB1 RPTOR WNT1 |
| 216 | GO:0042035 | regulation of cytokine biosynthetic process | 4.807E-01 | 2.43E+01 | 3 | IL10 IL12B NFkB1 |
| 216 | GO:0045927 | positive regulation of growth | 4.977E-01 | 2.504E+01 | 4 | ADAM10 ADAM17 MTOR RPTOR |
| 224 | GO:0002897 | regulation of immune effector process | 5.039E-01 | 2.524E+01 | 5 | CADM1 JL10 IL12B STAT1 STAT6 |
| 221 | GO:0042832 | defense response to protozoan | 5.154E-01 | 2.536E+01 | 2 | IL10 IL12B |
| 222 | GO:0005059 | membrane protein ectodomain proteolysis | 5.154E-01 | 2.536E+01 | 2 | ADAM10 ADAM17 |
| 222 | GO:0006359 | regulation of transcription by RNA polymerase III | 5.154E-01 | 2.536E+01 | 2 | MTOR RPTOR |
| 224 | GO:0031324 | negative regulation of cellular metabolic process | 5.154E-01 | 2.536E+01 | 11 | IL10 MTOR NFkB1 NFKB1 IRG3S3 RPTOR STAT1 STAT6 TNSL WNT1 ZC3H12D |
| 222 | GO:2000112 | regulation of cellular macromolecule biosynthetic process | 5.464E-01 | 2.673E+01 | 14 | CXCL10 FOXPA4 HCK IL10 MTOR NFAT5 NFkB1 NFKB1 IRG3S3 RPTOR STAT1 STAT6 TNSL WNT1 ZC3H12D |
| 222 | GO:0032269 | negative regulation of cellular protein metabolic process | 5.495E-01 | 2.673E+01 | 7 | IL10 MTOR NFkB1 IRG3S3 RPTOR WNT1 ZC3H12D |
| 221 | GO:0002285 | lymphocyte activation involved in immune response | 5.507E-01 | 2.673E+01 | 3 | IL12B MTOR STAT6 |
| 222 | GO:0002861 | regulation of inflammatory response to antigenic stimulus | 5.618E-01 | 2.703E+01 | 2 | IL10 IL12B |
| 224 | GO:0001582 | response to protozoan | 5.618E-01 | 2.703E+01 | 2 | IL10 IL12B |
| 224 | GO:0071375 | cellular response to peptide hormone stimulus | 5.751E-01 | 2.755E+01 | 4 | NFKB1 STAT1 STAT6 WNT1 |
| 231 | GO:0009888 | tissue development | 6.077E-01 | 2.873E+01 | 9 | ADAM17 HCK IL10 MTOR NPPY ROBO1 STAT1 STAT6 WNT1 |
| 221 | GO:0032107 | regulation of response to nutrient levels | 6.101E-01 | 2.873E+01 | 2 | MTOR NPPY |
| 234 | GO:0051220 | cytoplasmic sequestering of protein | 6.101E-01 | 2.873E+01 | 2 | IL10 TNSL |
| 234 | GO:0032104 | regulation of response to extracellular stimulus | 6.101E-01 | 2.873E+01 | 2 | MTOR NPPY |
| 234 | GO:0031323 | regulation of cellular metabolic process | 6.29E-01 | 2.949E+01 | 18 | ADAM17 CXCL10 FOXPA4 HCK IL10 IL12B MTOR NFAT5 NFkB1 NFKB1 IRG3S3 ROBO1 RPTOR STAT1 STAT6 TNSL WNT1 ZC3H12D |
| 234 | GO:0007162 | negative regulation of cell adhesion | 6.342E-01 | 2.961E+01 | 4 | ADAM10 IL10 PLXNB1 WNT1 |
| 231 | GO:0006355 | regulation of transcription, DNA-templated | 6.451E-01 | 2.999E+01 | 13 | CXCL10 FOXPA4 HCK IL10 MTOR NFAT5 NFkB1 NFKB1 IRG3S3 RPTOR STAT1 STAT6 TNSL WNT1 |
| 234 | GO:0043368 | positive T cell selection | 6.603E-01 | 3.032E+01 | 2 | MTOR STAT6 |
| 234 | GO:0002230 | positive regulation of defense response to virus by host | 6.603E-01 | 3.032E+01 | 2 | IL12B STAT1 |
| 244 | GO:1903203 | regulation of oxidative stress-induced neuron death | 6.603E-01 | 3.032E+01 | 2 | IL10 WNT1 |
| 241 | GO:0032258 | regulation of cellular protein metabolic process | 6.765E-01 | 3.083E+01 | 11 | ADAM17 CXCL10 IL10 IL12B MTOR NFkB1 IRG3S3 ROBO1 RPTOR WNT1 ZC3H12D |
| 244 | GO:0031347 | regulation of defense response | 6.794E-01 | 3.084E+01 | 6 | CADM1 HCK IL10 IL12B NFkB1 STAT1 |
| 244 | GO:0030335 | positive regulation of cell migration | 6.831E-01 | 3.086E+01 | 5 | ADAM10 ADAM17 CXCL10 IL8 MTOR |
| 244 | GO:0043900 | regulation of multi-organism process | 7.23E-01 | 3.265E+01 | 6 | CADM1 HCK IL12B IL8 NFkB1 STAT1 |
| 244 | GO:0045765 | regulation of angiogenesis | 7.453E-01 | 3.352E+01 | 4 | CXCL10 IL10 IL8 STAT1 |
| 244 | GO:1903506 | regulation of nucleic acid-templated transcription | 7.594E-01 | 3.392E+01 | 13 | CXCL10 FOXPA4 HCK IL10 MTOR NFAT5 NFkB1 NFKB1 IRG3S3 RPTOR STAT1 STAT6 TNSL WNT1 |
| 244 | GO:0019219 | regulation of nucleobase-containing compound metabolic process | 7.637E-01 | 3.392E+01 | 14 | CXCL10 FOXPA4 HCK IL10 MTOR NFAT5 NFkB1 NFKB1 IRG3S3 RPTOR STAT1 STAT6 TNSL WNT1 ZC3H12D |
| 244 | GO:1902745 | positive regulation of lamellipodium organization | 7.666E-01 | 3.392E+01 | 2 | MTOR WNT1 |
| 244 | GO:0031929 | TOR signaling | 7.666E-01 | 3.392E+01 | 2 | MTOR RPTOR |
| 254 | GO:2001141 | regulation of RNA biosynthetic process | 7.7E-01 | 3.394E+01 | 13 | CXCL10 FOXPA4 HCK IL10 MTOR NFAT5 NFkB1 NFKB1 IRG3S3 RPTOR STAT1 STAT6 TNSL WNT1 |
| 251 | GO:0001932 | regulation of protein phosphorylation | 7.748E-01 | 3.402E+01 | 8 | ADAM17 CXCL10 IL12B MTOR IRG3S3 ROBO1 RPTOR WNT1 |
| 254 | GO:0051248 | negative regulation of protein metabolic process | 7.984E-01 | 3.491E+01 | 7 | IL10 MTOR NFkB1 IRG3S3 RPTOR WNT1 ZC3H12D |
| 254 | GO:0007219 | Notch signaling pathway | 8.178E-01 | 3.555E+01 | 3 | ADAM10 ADAM17 STAT1 |
| 251 | GO:0032647 | regulation of interferon-alpha production | 8.226E-01 | 3.555E+01 | 2 | IL10 STAT1 |
| 254 | GO:0045954 | positive regulation of natural killer cell mediated cytotoxicity | 8.226E-01 | 3.555E+01 | 2 | CADM1 JL12B |
| 254 | GO:2000147 | positive regulation of cell motility | 8.317E-01 | 3.58E+01 | 5 | ADAM10 ADAM17 CXCL10 IL8 MTOR |
| 251 | GO:0002831 | regulation of response to biotic stimulus | 8.39E-01 | 3.597E+01 | 5 | CADM1 HCK IL12B NFkB1 STAT1 |
| 254 | GO:0010591 | regulation of lamellipodium assembly | 8.806E-01 | 3.761E+01 | 2 | MTOR WNT1 |
| 254 | GO:0002705 | positive regulation of leukocyte mediated immunity | 9.356E-01 | 3.955E+01 | 3 | CADM1 JL12B STAT6 |
| 254 | GO:0045860 | positive regulation of protein kinase activity | 9.381E-01 | 3.955E+01 | 5 | ADAM17 IL12B MTOR ROBO1 RPTOR |
| 261 | GO:0042036 | negative regulation of cytokine biosynthetic process | 9.405E-01 | 3.955E+01 | 2 | IL10 NFkB1 |
| 261 | GO:1902895 | positive regulation of pr-miRNA transcription by RNA polymerase III | 9.405E-01 | 3.955E+01 | 2 | IL10 NFKB1 |
| 264 | GO:0051272 | positive regulation of cellular component movement | 9.46E-01 | 3.964E+01 | 5 | ADAM10 ADAM17 CXCL10 IL8 MTOR |
| 264 | GO:0071407 | cellular response to organic cyclic compound | 9.866E-01 | 4.118E+01 | 5 | IL10 NFkB1 NFKB1 IRG3S3 SLC6A4 STAT1 |
| 264 | GO:0002717 | positive regulation of natural killer cell mediated immunity | 1.002E+00 | 4.152E+01 | 2 | CADM1 JL12B |
| 264 | GO:0071354 | cellular response to interleukin-6 | 1.002E+00 | 4.152E+01 | 2 | NFKB1 STAT1 |
| 264 | GO:0050728 | negative regulation of inflammatory response | 1.02E+00 | 4.209E+01 | 3 | IL10 IL12B NFkB1 |
| 264 | GO:0050776 | regulation of immune response | 1.046E+00 | 4.303E+01 | 7 | CADM1 HCK IL10 IL12B NFkB1 STAT1 STAT6 |
| 264 | GO:0006935 | chemotaxis | 1.054E+00 | 4.319E+01 | 5 | CXCL10 IL10 IL8 PLXNB1 ROBO1 |
| 274 | GO:0045191 | regulation of isotype switching | 1.066E+00 | 4.342E+01 | 2 | IL10 STAT6 |
| 274 | GO:0042330 | taxis | 1.072E+00 | 4.342E+01 | 5 | CXCL10 IL10 IL8 PLXNB1 ROBO1 |
| 274 | GO:0040017 | positive regulation of locomotion | 1.072E+00 | 4.342E+01 | 5 | ADAM10 ADAM17 CXCL10 IL8 MTOR |
| 274 | GO:0106056 | regulation of calcineurin-mediated signaling | 1.132E+00 | 4.55E+01 | 2 | MTOR NFAT5 |
| 274 | GO:0070884 | regulation of calcineurin-NFAT signaling cascade | 1.132E+00 | 4.55E+01 | 2 | MTOR NFAT5 |
| 274 | GO:0051246 | regulation of protein metabolic process | 1.161E+00 | 4.65E+01 | 11 | ADAM17 CXCL10 IL10 IL12B MTOR NFkB1 IRG3S3 ROBO1 RPTOR WNT1 ZC3H12D |
| 274 | GO:0030595 | leukocyte chemotaxis | 1.203E+00 | 4.784E+01 | 3 | CXCL10 IL10 IL8 |
| 274 | GO:0002822 | regulation of adaptive immune response based on somatic recombination | 1.203E+00 | 4.784E+01 | 3 | IL10 IL12B STAT6 |
| 274 | GO:1901653 | cellular response to peptide | 1.216E+00 | 4.819E+01 | 4 | NFKB1 STAT1 STAT6 WNT1 |
| 274 | GO:0051726 | regulation of cell cycle | 1.231E+00 | 4.862E+01 | 7 | ADAM17 IL10 IL12B IL8 MTOR RPTOR SLC6A4 |
| 284 | GO:0048856 | anatomical structure development | 1.268E+00 | 4.956E+01 | 16 | ADAM10 ADAM17 CADM1 CXCL10 HCK IL10 IL12B IL8 MTOR NPPY PLXNB1 ROBO1 SLC6A4 STAT1 STAT6 WNT1 |
| 281 | GO:0070741 | response to interleukin-6 | 1.268E+00 | 4.956E+01 | 2 | NFKB1 STAT1 |
| 284 | GO:1900087 | positive regulation of G1/S transition of mitotic cell cycle | 1.268E+00 | 4.956E+01 | 2 | ADAM17 RPTOR |
| 284 | GO:0050777 | negative regulation of immune response | 1.276E+00 | 4.969E+01 | 3 | IL10 IL12B STAT6 |