

cellular response to lipopolysaccharide, GO:0071222

- transcriptional repressor activity, GO:0001078
- interferon-gamma-mediated signaling pathway, GO:0060333
- response to virus, GO:0009615
- type I interferon signaling pathway, GO:0060337
- tumor necrosis factor receptor binding, GO:0005164
- positive regulation of B cell activation, GO:0050871
- NF-kappaB binding, GO:0051059
- steroid hormone mediated signaling pathway, GO:0043401
- negative regulation of T cell proliferation, GO:0042130
- steroid hormone receptor activity, GO:0003707
- regulation of sequence-specific DNA binding transcription factor activity, GO:0051090
- activation of NF-kappaB-inducing kinase activity, GO:0007250
- RNA polymerase II transcription factor activity, GO:0004879
- cellular defense response, GO:0006968
- positive regulation of interferon-gamma production, GO:0032729
- positive regulation of interleukin-2 production, GO:0032743
- cytokine receptor activity, GO:0004896
- tumor necrosis factor-activated receptor activity, GO:0005031
- regulatory region DNA binding, GO:0000975
- cellular response to lipoteichoic acid, GO:0071223
- regulation of cytokine secretion, GO:0050707
- transcriptional activator activity, GO:0001205
- response to muramyl dipeptide, GO:0032495
- phosphate ion binding, GO:0042301
- intrinsic component of plasma membrane, GO:0031226
- T cell proliferation, GO:0042098
- cytokine-mediated signaling pathway, GO:0019221
- 3', GO:0004114
- cellular response to estrogen stimulus, GO:0071391
- monocyte chemotaxis, GO:0002548
- toll-like receptor 9 signaling pathway, GO:0034162
- cellular response to cytokine stimulus, GO:0071345
- negative regulation of interferon-gamma production, GO:0032689
- platelet dense granule organization, GO:0060155
- retinoic acid binding, GO:0001972
- response to exogenous dsRNA, GO:0043330
- natural killer cell differentiation, GO:0001779
- positive regulation of alpha-beta T cell proliferation, GO:0046641
- positive regulation of natural killer cell proliferation, GO:0032819
- positive regulation of cytokine production, GO:0001819
- toll-like receptor 4 signaling pathway, GO:0034142
- serine-type endopeptidase inhibitor activity, GO:0004867
- response to peptide, GO:1901652
- ATP hydrolysis coupled anion transmembrane transport, GO:0099133
- positive regulation of interleukin-10 production, GO:0032733
- pentose-phosphate shunt, GO:0009052
- cellular response to gonadotropin stimulus, GO:0071371
- positive regulation of T-helper 2 cell differentiation, GO:0045630
- response to pain, GO:0048265
- cytidine deamination, GO:0009972
- positive regulation of interferon-gamma biosynthetic process, GO:0045078
- estrogen biosynthetic process, GO:0006703
- transmembrane receptor protein tyrosine kinase adaptor activity, GO:0005068
- interleukin-1 beta secretion, GO:0050702
- thyroid hormone receptor activity, GO:0004887
- negative regulation of myoblast differentiation, GO:0045662