

positive regulation of I-kappaB kinase/NF-kappaB signaling, GO:0043123

- protein autophosphorylation, GO:0046777
- ubiquitin ligase complex, GO:0000151
- positive regulation of NF-kappaB transcription factor activity, GO:0051092
- response to lipopolysaccharide, GO:0032496
- activation of cysteine-type endopeptidase activity involved in apoptotic process, GO:0006919
- cellular response to lipopolysaccharide, GO:0071222
- I-kappaB kinase/NF-kappaB signaling, GO:0007249
- regulation of mitotic cell cycle, GO:0007346
- defense response to virus, GO:0051607
- positive regulation of angiogenesis, GO:0045766
- receptor complex, GO:0043235
- activation of protein kinase activity, GO:0032147
- interferon-gamma-mediated signaling pathway, GO:0060333
- positive regulation of B cell activation, GO:0050871
- apoptotic signaling pathway, GO:0097190
- response to virus, GO:0009615
- ossification, GO:0001503
- JNK cascade, GO:0007254
- cellular response to organic cyclic compound, GO:0071407
- activation of MAPK activity, GO:0000187
- endoplasmic reticulum quality control compartment, GO:0044322
- regulation of defense response to virus by virus, GO:0050690
- extrinsic apoptotic signaling pathway via death domain receptors, GO:0008625
- activation of MAPKK activity, GO:0000186
- Hsp90 protein binding, GO:0051879
- Hsp70 protein binding, GO:0030544
- cytoplasmic side of plasma membrane, GO:0009898
- Notch signaling pathway, GO:0007219
- negative regulation of viral genome replication, GO:0045071
- T cell activation, GO:0042110
- Rho GTPase binding, GO:0017048
- negative regulation of transforming growth factor beta receptor signaling pathway, GO:0030512
- positive regulation of T cell proliferation, GO:0042102
- negative regulation of extrinsic apoptotic signaling pathway, GO:2001237
- NF-kappaB binding, GO:0051059
- positive regulation of extrinsic apoptotic signaling pathway, GO:2001238
- extrinsic apoptotic signaling pathway, GO:0097191
- extrinsic apoptotic signaling pathway in absence of ligand, GO:0097192
- tumor necrosis factor receptor binding, GO:0005164
- type I interferon signaling pathway, GO:0060337
- SMAD binding, GO:0046332
- cellular response to drug, GO:0035690
- positive regulation of cell death, GO:0010942
- positive regulation of smooth muscle cell proliferation, GO:0048661
- positive regulation of interferon-gamma production, GO:0032729
- T cell proliferation, GO:0042098
- cytokine-mediated signaling pathway, GO:0019221
- stress-activated protein kinase signaling cascade, GO:0031098
- response to interleukin-1, GO:0070555
- negative regulation of tumor necrosis factor production, GO:0032720
- release of sequestered calcium ion into cytosol, GO:0051209
- cellular iron ion homeostasis, GO:0006879
- positive regulation of cell division, GO:0051781
- epithelial to mesenchymal transition, GO:0001837
- multivesicular body, GO:0005771
- positive regulation of tumor necrosis factor production, GO:0032760
- negative regulation of MAP kinase activity, GO:0043407
- positive regulation of inflammatory response, GO:0050729
- negative regulation of viral transcription, GO:0032897
- amyloid-beta binding, GO:0001540
- spleen development, GO:0048536
- ubiquitin-ubiquitin ligase activity, GO:0034450
- positive regulation of defense response to virus by host, GO:0002230
- positive regulation of neuron differentiation, GO:0045666
- apoptotic cell clearance, GO:0043277
- positive regulation of stress-activated MAPK cascade, GO:0032874
- mitochondrial fission, GO:0000266
- positive regulation of phosphatidylinositol 3-kinase signaling, GO:0014068
- regulation of proteasomal protein catabolic process, GO:0061136
- positive regulation of protein export from nucleus, GO:0046827
- regulation of innate immune response, GO:0045088
- regulation of extrinsic apoptotic signaling pathway via death domain receptors, GO:1902041
- anoikis, GO:0043276
- collagen trimer, GO:0005581
- intrinsic component of plasma membrane, GO:0031226
- protein-cysteine S-palmitoyltransferase activity, GO:0019706
- response to ATP, GO:0033198
- positive regulation of proteolysis, GO:0045862
- positive regulation of transcription factor import into nucleus, GO:0042993
- dendritic cell chemotaxis, GO:0002407
- tumor necrosis factor-activated receptor activity, GO:0005031
- positive regulation of vascular endothelial growth factor receptor signaling pathway, GO:0030949
- hair follicle development, GO:0001942
- positive regulation of viral entry into host cell, GO:0046598
- transcriptional activator activity, GO:0001205
- positive regulation of interleukin-6 production, GO:0032755
- negative regulation of extrinsic apoptotic signaling pathway via death domain receptors, GO:1902042
- cellular response to amyloid-beta, GO:1904646
- positive regulation of cytokine production, GO:0001819
- positive regulation of protein homooligomerization, GO:0032464
- cellular response to interferon-gamma, GO:0071346
- interleukin-1-mediated signaling pathway, GO:0070498
- negative regulation of inflammatory response, GO:0050728
- negative regulation of insulin receptor signaling pathway, GO:0046627
- MAP kinase kinase kinase activity, GO:0004709
- positive regulation of cytosolic calcium ion concentration involved in phospholipase C-activating G-protein coupled signaling pathway, GO:0051482
- positive regulation of blood vessel diameter, GO:0097755
- response to tumor necrosis factor, GO:0034612
- positive regulation of Rho protein signal transduction, GO:0035025
- myeloid dendritic cell differentiation, GO:0043011
- response to interferon-alpha, GO:0035455
- cellular response to exogenous dsRNA, GO:0071360
- positive regulation of protein complex assembly, GO:0031334
- activin binding, GO:0048185
- defense response, GO:0006952
- positive regulation of B cell proliferation, GO:0030890
- metal ion transport, GO:0030001
- anterior/posterior axis specification, GO:0009948
- cardiac muscle hypertrophy, GO:0003300
- activation of NF-kappaB-inducing kinase activity, GO:0007250
- type I transforming growth factor beta receptor binding, GO:0034713
- response to interferon-beta, GO:0035456
- cytokine secretion, GO:0050663
- phospholipase C-activating G-protein coupled receptor signaling pathway, GO:0007200
- negative regulation of DNA biosynthetic process, GO:2000279
- negative regulation of interferon-gamma production, GO:0032689
- positive regulation of tumor necrosis factor secretion, GO:1904469
- negative regulation of interleukin-6 production, GO:0032715
- positive regulation of cell motility, GO:2000147
- midbrain development, GO:0030901
- positive regulation of ubiquitin-protein transferase activity, GO:0051443
- negative regulation of sodium ion transmembrane transporter activity, GO:2000650