R-HSA-1280215: Cytokine Signaling in Immune system
GO:0009615: response to virus
GO:0031347: regulation of defense response
GO:0006954: inflammatory response
hsa04668: TNF signaling pathway
GO:0002683: negative regulation of immune system process
GO:0030155: regulation of cell adhesion
GO:0035239: tube morphogenesis
WP5115: Network map of SARS-CoV-2 signaling pathway
GO:0030335: positive regulation of cell migration
GO:0008285: negative regulation of cell population proliferation
WP5088: Prostaglandin signaling
GO:0048568: embryonic organ development
hsa05200: Pathways in cancer
GO:0034612: response to tumor necrosis factor
GO:0001775: cell activation
GO:0006935: chemotaxis
GO:0043122: regulation of I-kappaB kinase/NF-kappaB signaling
GO:0045596: negative regulation of cell differentiation
hsa04621: NOD-like receptor signaling pathway
GO:0043410: positive regulation of MAPK cascade
hsa05164: Influenza A
WP5087: Malignant pleural mesothelioma
WP5039: SARS-CoV-2 innate immunity evasion and cell-specific immune response
GO:0001501: skeletal system development
GO:0050778: positive regulation of immune response
WP5094: Orexin receptor pathway
GO:0052548: regulation of endopeptidase activity
R-HSA-372790: Signaling by GPCR
R-HSA-6785807: Interleukin-4 and Interleukin-13 signaling
GO:0098586: cellular response to virus
GO:0050731: positive regulation of peptidyl-tyrosine phosphorylation
GO:0001934: positive regulation of protein phosphorylation
GO:0048608: reproductive structure development
GO:0061061: muscle structure development
WP619: Type II interferon signaling (IFNG)
GO:0043086: negative regulation of catalytic activity
GO:0048729: tissue morphogenesis
GO:0002697: regulation of immune effector process
GO:0009725: response to hormone
GO:0050866: negative regulation of cell activation
GO:0035456: response to interferon-beta
WP254: Apoptosis
GO:0007507: heart development
GO:0120035: regulation of plasma membrane bounded cell projection organization
GO:0002221: pattern recognition receptor signaling pathway
GO:1902532: negative regulation of intracellular signal transduction
hsa04217: Necroptosis
GO:0060284: regulation of cell development
GO:0003013: circulatory system process
GO:0051345: positive regulation of hydrolase activity
GO:0010632: regulation of epithelial cell migration
GO:0032755: positive regulation of interleukin-6 production
WP4655: Cytosolic DNA-sensing pathway
GO:0001667: ameboidal-type cell migration
GO:0040008: regulation of growth
GO:0001666: response to hypoxia
GO:0030856: regulation of epithelial cell differentiation
WP3624: Lung fibrosis
GO:0010942: positive regulation of cell death
GO:0045637: regulation of myeloid cell differentiation
GO:0048660: regulation of smooth muscle cell proliferation
R-HSA-1474244: Extracellular matrix organization
GO:2000027: regulation of animal organ morphogenesis
hsa05202: Transcriptional misregulation in cancer
WP2864: Apoptosis-related network due to altered Notch3 in ovarian cancer
GO:0050678: regulation of epithelial cell proliferation
GO:0001655: urogenital system development
GO:0032649: regulation of interferon-gamma production
GO:0051147: regulation of muscle cell differentiation
GO:0050878: regulation of body fluid levels
GO:0051051: negative regulation of transport
GO:0051046: regulation of secretion
GO:2001236: regulation of extrinsic apoptotic signaling pathway
hsa04933: AGE-RAGE signaling pathway in diabetic complications
GO:0048863: stem cell differentiation
GO:0002274: myeloid leukocyte activation
GO:0040013: negative regulation of locomotion
WP2880: Glucocorticoid receptor pathway
GO:0060760: positive regulation of response to cytokine stimulus
GO:0080135: regulation of cellular response to stress
GO:0044706: multi-multicellular organism process
hsa04625: C-type lectin receptor signaling pathway
GO:0043269: regulation of ion transport
GO:0001933: negative regulation of protein phosphorylation
WP2064: Neural crest differentiation
GO:0050691: regulation of defense response to virus by host
GO:0009612: response to mechanical stimulus
GO:0050819: negative regulation of coagulation
GO:0071496: cellular response to external stimulus
GO:0046634: regulation of alpha-beta T cell activation
GO:1903707: negative regulation of hemopoiesis
M196: PID IL23 PATHWAY
hsa04010: MAPK signaling pathway
WP3888: VEGFA-VEGFR2 signaling pathway
R-HSA-109582: Hemostasis
GO:0060322: head development
GO:0060021: roof of mouth development
R-HSA-1280218: Adaptive Immune System
GO:0030099: myeloid cell differentiation