



ID	Description
GO:0009615	response to virus
GO:0051607	defense response to virus
GO:0140546	defense response to symbiont
GO:0060337	type I interferon signaling pathway
GO:0071357	cellular response to type I interferon
GO:0034340	response to type I interferon
GO:0034341	response to interferon–gamma
GO:0048525	negative regulation of viral process
GO:0071346	cellular response to interferon–gamma
GO:0009991	response to extracellular stimulus
GO:0045071	negative regulation of viral genome replication
GO:0060333	interferon–gamma–mediated signaling pathway
GO:0031667	response to nutrient levels
GO:0002831	regulation of response to biotic stimulus
GO:0050792	regulation of viral process
GO:0019058	viral life cycle
GO:0043903	regulation of biological process involved in symbiotic interaction
GO:0045069	regulation of viral genome replication
GO:1903900	regulation of viral life cycle
GO:0071347	cellular response to interleukin–1
GO:0070555	response to interleukin–1
GO:0019079	viral genome replication
GO:0060759	regulation of response to cytokine stimulus
GO:0071496	cellular response to external stimulus
GO:0002683	negative regulation of immune system process
GO:0032496	response to lipopolysaccharide
GO:0071216	cellular response to biotic stimulus
GO:0071222	cellular response to lipopolysaccharide
GO:0001959	regulation of cytokine–mediated signaling pathway
GO:0002237	response to molecule of bacterial origin
GO:0016125	sterol metabolic process
GO:0071219	cellular response to molecule of bacterial origin
GO:0031669	cellular response to nutrient levels
GO:0031668	cellular response to extracellular stimulus
GO:0032479	regulation of type I interferon production
GO:0032606	type I interferon production
GO:0008203	cholesterol metabolic process
GO:0045088	regulation of innate immune response
GO:0001819	positive regulation of cytokine production
GO:1902652	secondary alcohol metabolic process
GO:0051098	regulation of binding
GO:0051090	regulation of DNA–binding transcription factor activity
GO:0002221	pattern recognition receptor signaling pathway
GO:0035456	response to interferon–beta
GO:0006695	cholesterol biosynthetic process
GO:1902653	secondary alcohol biosynthetic process
GO:0019216	regulation of lipid metabolic process
GO:0051235	maintenance of location
GO:0045444	fat cell differentiation
GO:0016126	sterol biosynthetic process
GO:0036293	response to decreased oxygen levels
GO:0002224	toll–like receptor signaling pathway
GO:0046890	regulation of lipid biosynthetic process
GO:0097193	intrinsic apoptotic signaling pathway
GO:0001818	negative regulation of cytokine production
GO:0045824	negative regulation of innate immune response
GO:0001666	response to hypoxia
GO:0070663	regulation of leukocyte proliferation
GO:0070482	response to oxygen levels
GO:0042594	response to starvation
GO:0048872	homeostasis of number of cells
GO:0002753	cytoplasmic pattern recognition receptor signaling pathway
GO:0060760	positive regulation of response to cytokine stimulus
GO:0050810	regulation of steroid biosynthetic process
GO:0009267	cellular response to starvation
GO:0051100	negative regulation of binding
GO:0062012	regulation of small molecule metabolic process
GO:0070059	intrinsic apoptotic signaling pathway in response to endoplasmic reticulum stress
GO:0051101	regulation of DNA binding
GO:0046718	viral entry into host cell
GO:0035455	response to interferon–alpha
GO:0051091	positive regulation of DNA–binding transcription factor activity
GO:0055094	response to lipoprotein particle
GO:0007584	response to nutrient
GO:0032480	negative regulation of type I interferon production
GO:0002832	negative regulation of response to biotic stimulus
GO:0045540	regulation of cholesterol biosynthetic process
GO:0106118	regulation of sterol biosynthetic process
GO:0071402	cellular response to lipoprotein particle stimulus
GO:0008202	steroid metabolic process
GO:0042493	response to drug
GO:0002262	myeloid cell homeostasis
GO:0070498	interleukin–1–mediated signaling pathway
GO:0019915	lipid storage
GO:0050727	regulation of inflammatory response
GO:0036003	positive regulation of transcription from RNA polymerase II promoter in response to stress
GO:0007249	I–kappaB kinase/NF–kappaB signaling
GO:0070661	leukocyte proliferation
GO:0043392	negative regulation of DNA binding
GO:0044409	entry into host
GO:0006984	ER–nucleus signaling pathway
GO:0002685	regulation of leukocyte migration
GO:0002687	positive regulation of leukocyte migration
GO:0050729	positive regulation of inflammatory response
GO:0043331	response to dsRNA
GO:0031331	positive regulation of cellular catabolic process
GO:0045598	regulation of fat cell differentiation
GO:0070423	nucleotide–binding oligomerization domain containing signaling pathway
GO:0071356	cellular response to tumor necrosis factor
GO:0043620	regulation of DNA–templated transcription in response to stress
GO:0001889	liver development
GO:0035872	nucleotide–binding domain, leucine rich repeat containing receptor signaling pathway
GO:1902930	regulation of alcohol biosynthetic process
GO:0031349	positive regulation of defense response
GO:0061008	hepaticobiliary system development
GO:0097191	extrinsic apoptotic signaling pathway
GO:0051701	biological process involved in interaction with host
GO:1902041	regulation of extrinsic apoptotic signaling pathway via death domain receptors
GO:0002697	regulation of immune effector process
GO:0032722	positive regulation of chemokine production
GO:1990830	cellular response to leukemia inhibitory factor
GO:0019218	regulation of steroid metabolic process
GO:0090181	regulation of cholesterol metabolic process
GO:1990823	response to leukemia inhibitory factor
GO:0052126	movement in host environment
GO:0062197	cellular response to chemical stress
GO:0050777	negative regulation of immune response
GO:0050873	brown fat cell differentiation
GO:2001233	regulation of apoptotic signaling pathway
GO:0034612	response to tumor necrosis factor
GO:0045600	positive regulation of fat cell differentiation
GO:0046165	alcohol biosynthetic process
GO:0001961	positive regulation of cytokine–mediated signaling pathway
GO:0043618	regulation of transcription from RNA polymerase II promoter in response to stress
GO:0070664	negative regulation of leukocyte proliferation
GO:0034599	cellular response to oxidative stress
GO:0051092	positive regulation of NF–kappaB transcription factor activity
GO:0043619	regulation of transcription from RNA polymerase II promoter in response to oxidative stress
GO:1990440	positive regulation of transcription from RNA polymerase II promoter in response to endoplasmic reticulum stress
GO:0036294	cellular response to decreased oxygen levels
GO:0043122	regulation of I–kappaB kinase/NF–kappaB signaling
GO:0003725	double–stranded RNA binding
GO:0005126	cytokine receptor binding