			regulation of MHC class II biosynthetic process (GO:0045346)
			regulation of intrinsic apoptotic signaling pathway in response to DNA damage by p53 class mediator (GO:1902165)
			regulation of DNA methylation (GO:0044030)
			negative regulation of protein acetylation (GO:1901984)
			negative regulation of peptidyl–lysine acetylation (GO:2000757)
			negative regulation of myoblast differentiation (GO:0045662)
			negative regulation of intrinsic apoptotic signaling pathway in response to DNA damage by p53 class mediator (GO:1902166)
			negative regulation of intrinsic apoptotic signaling pathway by p53 class mediator (GO:1902254)
			negative regulation of histone acetylation (GO:0035067)
			macrophage differentiation (GO:0030225)
			lymphoid progenitor cell differentiation (GO:0002320)
			histone H2A ubiquitination (GO:0033522)
			anatomical structure regression (GO:0060033)
			granulocyte differentiation (GO:0030851)
			forebrain morphogenesis (GO:0048853)
			lateral ventricle development (GO:0021670)
			proteolysis involved in cellular protein catabolic process (GO:0051603)
			ubiquitin-dependent protein catabolic process (GO:0006511)
			modification-dependent macromolecule catabolic process (GO:0043632)
			modification-dependent protein catabolic process (GO:0019941)
			midbrain development (GO:0030901)
			protein deubiquitination (GO:0016579)
			protein modification by small protein removal (GO:0070646)
			negative regulation of peptidase activity (GO:0010466)
			DNA recombination (GO:0006310)
			transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0007178)
			cellular response to transforming growth factor beta stimulus (GO:0071560)
			response to transforming growth factor beta (GO:0071559)
			regulation of cellular response to growth factor stimulus (GO:0090287)
			regulation of transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0090092)
			negative regulation of transforming growth factor beta receptor signaling pathway (GO:0030512)
			regulation of transforming growth factor beta receptor signaling pathway (GO:0017015)
			negative regulation of cellular response to growth factor stimulus (GO:0090288)
			negative regulation of transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0090101)
			regulation of proteasomal protein catabolic process (GO:0061136)
			transforming growth factor beta receptor signaling pathway (GO:0007179)
			regulation of cellular protein catabolic process (GO:1903362)
			regulation of proteolysis involved in cellular protein catabolic process (GO:1903050)
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