			regulation of mammary gland epithelial cell proliferation (GO:0033599) positive regulation of receptor biosynthetic process (GO:0010870) positive regulation of muscle cell apoptotic process (GO:0010661)	
			centriole assembly (GO:0098534)	
			centriole replication (GO:0007099)	
			histone H2A ubiquitination (GO:0033522)	
			regulation of sulfur metabolic process (GO:0042762) regulation of cofactor metabolic process (GO:0051193)	
			regulation of coenzyme metabolic process (GO:0051196)	
			regulation of acyl-CoA biosynthetic process (GO:0050812)	
			regulation of acetyl–CoA biosynthetic process from pyruvate (GO:0010510)	
			positive regulation of potassium ion transport (GO:0043268)	
			negative regulation of metanephros development (GO:0072217)	
			histone ubiquitination (GO:0016574)	
			granulocyte differentiation (GO:0030851)	
			amino acid transmembrane transport (GO:0003333) face development (GO:0060324)	
			negative regulation of microtubule polymerization or depolymerization (GO:0031111)	
			tube morphogenesis (GO:0035239)	
			regulation of microtubule depolymerization (GO:0031114)	
			negative regulation of microtubule depolymerization (GO:0007026)	
			body morphogenesis (GO:0010171)	
			epithelial tube morphogenesis (GO:0060562)	
			cellular protein complex assembly (GO:0043623) regulation of peptidase activity (GO:0052547)	
			organ morphogenesis (GO:0009887)	
			pigmentation (GO:0043473)	
			intracellular receptor signaling pathway (GO:0030522)	
			chromatin modification (GO:0016568)	
			nitrogen compound transport (GO:0071705)	
			response to molecule of bacterial origin (GO:0002237)	
			regulation of peptide transport (GO:0090087)	
			response to lipopolysaccharide (GO:0032496) MyD88-dependent toll-like receptor signaling pathway (GO:0002755)	
			toll–like receptor TLR1:TLR2 signaling pathway (GO:0038123)	
			signal transduction by phosphorylation (GO:0023014)	
			MAPK cascade (GO:0000165)	
			negative regulation of cell-substrate adhesion (GO:0010812)	
			RNA splicing (GO:0008380)	
			regulation of skeletal muscle tissue development (GO:0048641)	
			negative regulation of organelle organization (GO:0010639) negative regulation of cell-matrix adhesion (GO:0001953)	
			ectodermal placode formation (GO:0060788)	
			brain development (GO:0007420)	
			central nervous system development (GO:0007417)	
			actin cytoskeleton organization (GO:0030036)	
			regulation of small GTPase mediated signal transduction (GO:0051056)	
			negative regulation of protein depolymerization (GO:1901880)	
			cognition (GO:0050890) negative regulation of protein complex disassembly (GO:0043242)	
			actin filament–based process (GO:0030029)	
			actin filament organization (GO:0007015)	
			carbohydrate derivative biosynthetic process (GO:1901137)	
			negative regulation of cytoskeleton organization (GO:0051494)	
			negative regulation of proteolysis (GO:0045861)	
			negative regulation of protein maturation (GO:1903318)	
_	_	_	negative regulation of protein processing (GO:0010955)	
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