

		0	10	20	30	40
	cellular component morphogenesis	GO:0032989				
	circulatory system development	GO:0032989				
	columnar/cuboidal epithelial cell development	GO:0032989				
	post-embryonic animal organ morphogenesis	GO:0032989				
	calcium ion homeostasis	GO:0032989				
	xenobiotic metabolic process	GO:0032989				
	male mating behavior	GO:0032989				
	pyruvate metabolic process	GO:0032989				
	nitrogen ion homeostasis	GO:0032989				
	metabolic anion transport	GO:0032989				
	motor neuron axon guidance	GO:0032989				
	purine-containing compound biosynthetic process	GO:0032989				
	regulation of cytokine catabolism of metalloprotease	GO:0032989				
	calcium-dependent cell-cell adhesion via plasma membrane cell adhesion molecules	GO:0032989				
	acid secretion	GO:0032989				
	cell-cell adhesion mediated by cadherin	GO:0032989				
	cell-cell adhesion mediated by integrin	GO:0032989				
	defense response to Gram-negative bacterium	GO:0032989				
	digestive system development	GO:0032989				
	microtubule-based transport	GO:0032989				
	neuroepithelial neural tube closure	GO:0032989				
	positive regulation of organ growth	GO:0032989				
	smoothed muscle cell differentiation	GO:0032989				
	auditory receptor cell morphogenesis	GO:0032989				
	embryonic cell morphogenesis	GO:0032989				
	establishment of imaginal disc-derived structure	GO:0032989				
	imaginal disc pattern formation	GO:0032989				
	negative regulation of organ growth	GO:0032989				
	protein autophosphorylation	GO:0032989				
	sensory perception of light stimulus	GO:0032989				
	glyoxylate metabolic process	GO:0032989				
	activation of cysteine-type endopeptidase activity involved in apoptosis	GO:0032989				
	chloride transmembrane transport	GO:0032989				
	lumen acidification	GO:0032989				
	glycosphingolipid metabolic process	GO:0032989				
	natural killer cell differentiation	GO:0032989				
	positive regulation of cytokine signaling	GO:0032989				
	positive regulation of cell growth	GO:0032989				
	protein NED-linked ubiquitination	GO:0032989				
	retroic, response to fatty acid	GO:0032989				
	steroid hormone mediated signaling pathway	GO:0032989				
	epithelial homeostasis	GO:0032989				
	activation of phosphatidylcholine 4-acyl	GO:0032989				
	activation of phospholipase A activity	GO:0032989				
	agranulocyte	GO:0032989				
	auditory receptor cell morphogenesis	GO:0032989				
	arachidonic acid metabolism	GO:0032989				
	cellular response to growth hormone stimulus	GO:0032989				
	central nervous system morphogenesis	GO:0032989				
	detection of mechanical stimulus	GO:0032989				
	detection of molecule of chemical origin	GO:0032989				
	establishment or maintenance of polarity of larval maggot disc epithelium	GO:0032989				
	facultative anaerobic respiration	GO:0032989				
	glutathione catabolism	GO:0032989				
	grooming behavior	GO:0032989				
	hemolymph	GO:0032989				
	long-chain fatty acid biosynthetic process	GO:0032989				
	mating behavior	GO:0032989				
	mitochondrial cytochrome c oxidase assembly	GO:0032989				
	monounsaturated fatty acid biosynthetic process	GO:0032989				
	negative regulation of acid vesicle morphogenesis	GO:0032989				
	negative regulation of cardiac muscle cell differentiation	GO:0032989				
	negative regulation of cardiac muscle tissue development	GO:0032989				
	negative regulation of histone gene expression	GO:0032989				
	negative regulation of steroid metabolic process	GO:0032989				
	phosphatidylinositol-3-phosphate biosynthetic process	GO:0032989				
	positive regulation of cholesterol esterification	GO:0032989				
	reduction of food intake in response to dietary excess	GO:0032989				
	regulation of sensory perception of pain	GO:0032989				
	response to pain	GO:0032989				
	salivary gland development	GO:0032989				
	unsaturated fatty acid metabolic process	GO:0032989				
	white fat cell differentiation	GO:0032989				
	alcohol catabolism	GO:0032989				
	cellular catabolic process	GO:0032989				
	clustering of voltage-gated calcium channels	GO:0032989				
	conductive acid balance function	GO:0032989				
	cuticle hydrolytic catabolic process	GO:0032989				
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