REVIGO Gene Ontology treemap

regulation of cellular process	positiv regulatio s biologio	n of regu	logical I	positive regulation of cellular	regulation of apoptotic	cellular component	tissue morphogenesis	anatomic structure formation involved	tissue	cell ent development	odontogenesis of dentin–containing tooth	embryo development	stem cel	on pos	'de novo' ttranslational otein folding ubiquitination	
contain process	proces	l pro	ocess	process	process	movement		morphogen					stem cell proliferation		primede novoprime posttranslational protein folding	
cell surface receptor signaling pathway	regulation o	regulati of cel	ll reg	gative ulation gnaling	negative egulation of cell	negative regulation of biological process	cardiovascular system development	morphogenesis of an epithelium		formation formation	of heart	prostate gland is development	mesenchyma proliferatio	al cell sr	protein diffication by nall protein dephosphorylation removal	
regulation of response	regulation of transcription from RNA polymerase II	negative regulation of response	T cell activation	actin filament-based process	regulation of macromolecul biosynthetic	negative regulation of chondrocyte	circulatory system development		of endothelial barrier	eproductive skir system development	ment developmer	tube morphogenesis	response cellular de	wounding efense	o molting cycle molting cycle	
negative regulation	promoter positive regulation	positive regulation	regulation of sequence–specifi	regulation of	process positive regulation	differentiation positive of regulation	nervous system	anterior/posterior axis specification	development	muscle structure evelopment	let proliferation	or lymphoid	respor response to hydrogen peroxide	wound healing	homeostasis glomerular	
	of phosphatidylinositol 3–kinase signaling	régulation de to stimulus	DNA binding of apoptotic factor activity	macromolecule process process		ve of metabolic process	development	axio opcomoduori		cell angioge orphogenesis involv involved in in wou	ed of a branching	· ·	develonmental		filtration	
signal transduction	glial cell	cell activatior	regulation of endopeptidas activity	regulation of multicellular organismal process	Intracellula	interleukin–1 production	skin morphogenesis	odontogenesis	myoblast er	ifferentiation healing cell cell prolifera	pattern specification	endoderm development	process	bi	ological regulation	
positive regulation of	regulation of biosynthetic	positive regulation of intracellular signal transduction	regulated secretory pathway	positive regulation of multicellular organismal process platelet-derived	negative regulation of multicellular organismal process regu	regulation of cell fate specification	· ·	single-multicellular organism process	ectoderm development	platelet notoch	enesis immune reg	julation of elopmental development	locomotion	cel prolifer	signaling	
process transcription from RNA	negative regulation	regulation of immune system process	positive regulation of regulation of	pathway	regulation of cell locali.	face	chemotaxis	response to external	response to respo	actomyosin	matrix	extracellular structure organization	response to stimulus	immur syster	organismal	
polymerase II promoter	of cytokine secretion	interleukin–1 beta production	tumor necrosis factor–mediated signaling pathway	morphogenesis re of a branching se	eceptor protein erine/threonine inase signaling of	ositive gulation cellular phosphate-mediated signaling		stimulus —chemotax	stimulus cytok	actomyos	in structure or oligomerization	cytoskeletal ganization at plasma		cellular	protein localization to	
cell communication	regulation of metabolic process	negative regulation of sequence—specific DNA binding transcription factor activity	enzyme linked receptor proteir signaling pathway	regulation of signaling	regulation mo	neutrophil ediated immunity	response to chemical	cellular response to stimulus	to	junction organization	cell junction organization		r	process involved in eproduction	cell surface	