REVIGO Gene Ontology treemap

						Gene Ontology	<u> </u>					
viral transcription	multi–organism metabolic proces			translational termination negative regulation of isotype switching to IgE isotypes	regulation of cell proliferation	regulation of calcium-dependent cell-cell adhesion	insulin receptor signaling pathway via phosphatidylinositol 3-kinase	positive regulation of Wnt signaling pathway, planar cell polarity pathway	response to cortisol	negative regulation of canonical Wnt signaling pathway	sinoatria	system I node pment
					negative regulation of calcium–dependent cell–cell adhesion	involved in ureter development negative regulation		signaling pathway 1	negative regulation of androgen receptor signaling pathway	neural plate pattern specification		
cellular macromolecule catabolic process	translational elongation	hypome of CpG	island is				regulation of cel	cell proliferation ureter development	regulation of eceptor recycling	negative regulation of anoikis		
membrane	oligosaccharide	cellular con	nponent el	transcription longation from NA polymerase	canonical Wnt signaling pathway involved in neura plate anterior/posterior pattern formation		response to retinoic acid	regulation of cell adhesion	process	enetic interleukin-4 printing production	microvillus microvillus assembly	microvillus assembly organization
organization	leucine metabolic			II promoter cyclooxygenase	regulation of genetic imprinting	positive regulation of integrin–mediated signaling pathway	fibroblast	or oubotrato	negative regulation of otch signaling	egative regulation of cellular process	cell proliferation	
ranslational initiation	process glutathione		multivesicular body sorting pathway	protein	to membrane	single–organism ntracellular transport sing	endocytosis	establishment of mitochondrion localization	clathrin-mediate endocytosis	ed pinocytosis	cellular iron ion	catabolic catabolis process
negative regulation of type 2 immune response	multi-organism biosynthetic process		clathrin coa assembly	t peptidyl-prolyl isomerization				positive regulation	import into	mitochondrion localization	homeostasis	is
chitin catabolic process	i interaction i	NA-templated transcription, elongation	autophagy	regulation of B cell mediated immunity	regulation of mitochondrial membrane permeability involved in programmed necrotic cell death	protein localization to endoplasmic reticulum	positive regulation of early endosome to late endosome transport	import into nucleus	transport	cellular localization	chronic inflammatory response	type 2 immune response