REVIGO Gene Ontology treemap

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neutrophil chemotaxis	taxis	leukocyte migration involved in inflammatory response	antigen processing and presentation of exogenous peptide antigen via MHC clas I, TAP-independent	of cell	presenta	ng and atrans enous me	sition etal Ca	cytosolic alcium ion transport response	extracellular matrix organization	cellular compone moveme	ent regu	lation of ocytosis	negative regulation of ion transmembrane transporter activity	response external stin	nulus respor	respons to woundir	response to ng chemical
response to glucocorticoid	regulation of response to external stimulus	response to nutrient levels	to metal oxy	response to oxygen–containing compound	growth factor receptor signaling pathway	transport	cation ransport	to inorganic substance	organization	protein heterotrimerization	regulation of cellular component		neuron recognition	regulation	response to	wound	inflammatory
		responsmeut extracellular	arachidonic rophilichen secretion	adenylate otaxis oparime receptor signaling pathway	to acid response	defense fresponse to fungus	extracellula regulation of signal transductio	gamma-delta T cell activation	extrac	nositive ellular matri regulation	positive x organiz	ation	positive regulation	to stimulu res	endogenous stimulus ponse to exte		response IS protein
response to organic cyclic compound	leukocyte migration	stimulus positive regulation	G-protein coupled	phospholipid efflux	negative regulation of lipid transpo	Communication	n by	response to carbohydrate	extracellular structure organization	of cytokine secretion positive	transport			positive regulation of response	response to stress	peptidyl-prolin- modification adaptive	peptidyl–proly isomerization
	response to	of apoptotic signaling pathway antigen processing and presentation	receptor protein signaling pathway	protein localization	response to fungus	second-messenger-med	dotosti	response to		regulation of ion transport	platelet degranulation negative	particle remodeling	organization g	to stimulus	complement activation	immune ac	rotein small molecule metabolic process
response to lipid	alcohol unsatura	of endogenous peptide antigen via MHC class I	of lipid transport	to extracellular region	lipid localizatio	proton of transport, down bacteriu		lipoprotein	hemidesmosome assembly	microvillus assembly	regulation of type I interferon production	egulation cell morphogenesis involved in differentiation cell cell morphogenesis er	negative regulation of transcription regulatory region DNA binding	inflammatory	lipid metabolic process	acute-phase response	detection of stimulus
single–multicellu organism proce	fatty ac biosynthe	id icosanoid biosynthetic process	Process	development	adrenal gland development m	ventricle	oulmonary artery rphogenesis	steroid esterification	positive mainte homeostatic	nance sys	of of	questering GFbeta in tracellular matrix		cellular	developmental	single–organisn process	locomotion
	cardiovaso system developm	fatty acid metabolic	organismal metabolic process	oositive regulation of nucleotide metabolic process	trabecula	derived foam	ung saccule evelopmen	accule regulation of phosphoprotein		action home	on eostasis	regulation of system	organism	al process			
collagen catabolic process		small molecticoll biosynthetic	agen catab catabolic	positive regulation of purine nucleotide olism regulation	s morphogene	nucleoside	regulation of axon regeneration	foam cell differentiation	of metal ion pos	itive seques	stering	regulation of body	glycerol-3-phosph metabolic proces glycerol-3	phosphate	behavior	response stimulus	regulation of multicellular organismal process
	organonitro compour biosynthe	ogen nd lipid	process single-organism biosynthetic	of JNK cascade cellular biogenic amine	production	of nucleotide metabolic proce	monocarbo	bolic		ation of structure excre		renal corption	metabolic process	coenzyme metabolic process	single–organis cellular proces		cell recognition
glycerol crea metabolic biosyn process proc	tine thetic trabecu	regulation of cholesterol	fatty acid derivative	metabolic process cellular aldehyde metabolic process	developmen process glucose metaboli process	metaboli process regulation of cAMP metab	S of coagula	positive	cell adhesio <mark>ce</mark>	neutr Il adhesion aggre	rophil adhes membra rophil gation	philic cell-cell ion via plasma ne cell adhesion nolecules ukocyte regation	biologica	l adhesion	regulation of growth	communication h	rganic ydroxy superoxide mpound metabolism