REVIGO Gene Ontology treemap

neutrophil chemotaxis	regulation of phagocytosis	anion transport	of ion transmem	egative regulation ion transmembrane ransporter activity		ntenance ocation	regulation of cellular component rganization	single-multice	ellular rena	al chemo	chemokine rei		primary alcohol catabolic	unsatura acid bios prod		i alueriyue	metabolic process	terpenoid catabolic process
	cellular component movement	regulation of ion transport	and presentation of	urea transport	organic acid transport	metal ion transport	ion transport	system process	interleukin–3	positive regulation granulocyte macropy colony–stimulating biosynthetic proc	phage factor ir	positive egulation of nterleukin–3 piosynthetic	icosanoid	proc	cess	ohol catab	process	autophagy
leukocyte migration	localization of cell	organic anion transport	chemical homeostasis	cellular phosphate ion homeostasis	glycoprotein transport	protein localization to juxtaparanode region of axon	tubulin complex assembly	regulation of nervous system	ion	e production ex	n xcretion	of cytokine	biosynthetic process	fatty deriv meta	ative abolic	regulation of cholesterol esterification	molecule biosynthetic process catal	
sequestering of metal ion	sodium ion transport		cyte migrat compound transport	cynanca	regulation of	negative regulation of sodium ion transport	efflux	development regulation of system	absorption cytokine production	system smooth muscle	muscle system	granulocyte macrophage colony–stimulating factor production	response alcohol	est	onse to	cellular response to mineralocorticoid stimulus		detection of endogenous stimulus
leukocyte migration involved in inflammatory	acid secretion	derivative	regulation of synapse organization	cellular proces	paranode region of axon regulation of	ion complex biogenes	positive regulation of sishomeostatic process	esophagus smooth muscle contraction	neuronal action potential	renal sodium ion in	negative regulation of nterferon-beta production	multicellular organismal signaling	response to	resp	onse to	detection of hormone stimulus	density lipopr particle	to tein response to transition metal nanoparticle
response sequestering of zinc ion	arachidonic acid secretion	glutathione transport	communication lipid localization	organizatio microvillus assembly	presentation	n secreting	microvillus organization	response to external stimulus	response to wounding	endopep	regulation of endopeptidase activity		DNA binding transcription factor activity	zymogen activation	neutrophil aggregation neutrophil aggr		leukocyte	multicellular organismal
positive regulation of apoptotic signaling pathway	positive regulation of T cell proliferation	positive regulation of cell communication	beta selection	myeloid cell activation involved in immune response	nuclear chang	apoptotic process	- 3 - 3	chronic inflammatory response	to stress	positive regulation of	and and a Con-	roomirotor	y compound metabolic process	pound transcript from RNA polymerase I promoter promoter		ukocyte ell adhesior	aggregation	process
adenylate cyclase–activating G–protein coupled receptor signaling	positive	dopamine receptor regulation of signaling pathway		neurotrophic factor receptor signaling pathway signaling pmineralocorticolor receptor signaling	clustering of voltage-gate sodium pathwaynels nuclear membran	signal transduction b phosphorylatio	of growth	serotonin secretion by platelet	s-activated lein kinase ignaling asscade regenerati	n	response to fungus		regulation of		regulation of multicellular organismal process		odontoblast differentiation odontoblast differentiation epidermis development	to stimulus
pathway regulation of ERK1 and ERK2 cascade	rogalation	G-protein coupled receptor signaling pathway, coupled to cyclic nucleotide second messenger	positive regulation of response to stimulus	pathway myeloid dendritic cell activation involved in immune response	organization leukocyte proliferation	singl	gnaling e e e e e e e e e e e e e e e e e e e	wound healing res	of positive regulation skeletal muscle tiss regeneration positive regulation skeletal muscle tiss positive regulation	defense to bac		se infla	ammatory sponse	negative regulation of macroautophagy	loco	motion	single–organism process	