## REVIGO Gene Ontology treemap

single-multicellula organism process		contraction	process	i tissue	linoleic a metabo proces	olic differen	li cell	nonocarboxylio acid biosynthetic process	response to external stimulu	response to	response	e to response to extracellula s stimulus	protein localization extracellu region	n to regulation	on of of lip	into host cell by promotion of host	regulation of lipoprotein lipase activity positive	CD24 biosynthetic process
	NAD metabolic process	cerebellar cortex development	aspartyl-tRNA aminoacylation	defecation		hypometh of CpG i	island p	insulin	response to wounding	response to stress	regulation of respons	se of response	modulation of phagocytosis in othe organism involved i symbiotic interactio	n extravagation	negative regulation o arachidonic acid secretio	regulation of entry of bacterium into	regulation of CD24 biosynthetic process	regulation of genetic imprinting
collagen catabolic process	bone	nervous system development	negative regulation of very-low-density lipoprotein particle clearance	embryo implantatior	regulation of type I	n regula I of bo	ody c	oagulation		ponse to ext	1 ( ) 1	ilusion collatera	protein le	endocytic ocalization t	a	on cohalami	regula lipopr lipase a	otein
creatine	positive regulation of myoblast	cell communication by electrical coupling involCO in cardiac conduction	peptide llagen cata amidation	cell morphogenesis <b>bolism</b> in differentiation	adrenal <sub>n</sub> gland development		protein peptidyl-proly isomerization	I of location	response to	positive regulation	reç	sprouting gulation esponse stress-induce	trans-Golgi network to recycling	endocytosis	ion regul	positive regulation of early endosome to late endosom	negative regulation of hydrolase activity negative	memory T cell activation
biosynthetic process	peptidyl-proline modification	fatty acid metabolic	peptide modification	complement activation, <sub>f</sub> classical pathway	nistone H3–T11 re phosphorylation	gulation of cell	negative regulation of low-density lipoprotein particle eceptor biosynthetic process	negative regulation of smoothened signaling pathway involved in dorsal/ventral neural tube patterning	chemical axon choice	to stimulus	positive reg	external premature senescence gulation	ocalization	phagocytosisb	ransport	transport  stein  sport  siderophore	regulation of phospholipase A2 activity positive	regulation of lipase activity  G2/M transition
negative regulation of heart rate	circulatory system	2-oxobutyrate biosynthetic process	primary amino compound biosynthetic	axon development	insulin metabolic process	eptidyl-pyroglutamic acid biosynthetic process, using glutaminyl-peptide cyclotransferase	positive egulation o	ol of	point recognition	cell migration	of T cell responded ox	ponse to kidative stress	or ceir	platelet degranulation		n lipid transport	regulation of memory T cell activation	of mitotic cell cycle
multicellular organismal metabolic process	development creatine metabolic process	activation of meiosis involved in egg activation	protein amidation	alcohol catabolic process	N-acetylneuraminate	positive regulation de	egulation on acrophage erived foan cell	organic hydroxy compound catabolic	extracellula	ar matriy	extracellular			dhesion  adhesion	negative of regulation of calcular-dependent of calcular-dependent or cal-cell subsession of cal-cell subsession of calcular dependent of calcular depende	gical adhesi		ry suckling r behavior ry behavior
response to	regulation of transforming growth factor beta receptor signaling pathway	cellular copper ion homeostasis	response to alcohol	cellular response to norepinephrine stimulus	response inorgani substand		oid c	cytosolic alcium ion transport	organiz			structure organization		of regulatio			short-term memory	reproductive behavior
	response to organic cyclic	cellular response to thyroid hormone stimulus	response to metal ion		l to l	response	terleukin–12–media signaling pathway	intracellular sequestering of iron ion	extracellular mat		rix organization		organization by cell–cell adhesion  antigen processing and presentation of		thesion mu essing org	lticellular ganismal	developmenta process	locomotion
of TGFbeta in extracellular matrix	insulin–like growth factor receptor	response to glu negative regulation of ryanodine-sensitive calcium-release channel activity response to carbohydra		response to azide		assimilation I by chelation and	ow-densi lipoproteii particle mediated signaling	n positive regulation of sequestering of calcium ior	hemidesmoso assembly	compo	nent trimerization organizatio		of endogeno antigen via Mipres	and procent	peptide pandiss endent	rocess	single–organism	
positive regulation of release of sequestered calcium ion into cytosol	response to transition metal nanoparticle	response to organic substance	cellular response to azide	positive regulation of ion transport	transmembrane receptor protein serine/threonine kinase signaling pathway	iron assimilation	sequestering of iron ion	1 reficulum	protein heterotrimeriza	cellu compo disasse	onent cytoskelet organizati	matrix disassemb	arantigen and prese of endog antig	enous preser peptide	sing and tation of antigen	gulation f signal	process single-organ cellular proc	