## REVIGO Gene Ontology treemap

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response to chemical	cellular response to stimulus	response to acid chemical	response to drug	response to biotic stimulus	respo	ipid	drug ansmembrane transport response to	vascular process in circulatory system	circulato system proces	multi-multice organism pr		brain egmentation	central nervous system segmentation	regulation of caveolin–mediated endocytosis	carnitine transmembrane transport	lipid localizatio	localization of cell	regulation of receptor-mediated endocytosis
chemotaxis	drug transport	response to other organism	response to oxygen–containing compound	response to anoxia	immune	to osmotic stress	decreased oxygen levels	rhombomere 5	rhombomer		re	egative egulation vascular	muscle contraction	caveolin-mediated endocytosis regulation	organic aci transport of caveolir	structui	e L. Golgi	negative regulation of integrin cytosis
response to	fatty acid	cellular response to response to chemical	response external stir	response to	regulation		of epithelial cell	development	cell	p		rmeability	interspecies	monocarboxylic acid transport	receptor-mediate endocytosis	nitric o	oort deriv	acid epithelial ative fluid
external stimu xenobiotic	Jius ,	stimulus response to	response to organic cyclic compound	response to stress	response		response to	regulation of system process		atory system	_		ent quorum sensing	lipid transport	extracellula matrix organizatio	sodium-de organic	pendent thyro	pid hormone
metabolic proc		oleic acid	chemokinesis	response to mineralocorticol	blood coagulatio extrinsion	on, c	ne	•	spongiotrophoblast layer development	angiogenesis de	cartilage evelopmen		n process	audini au	hepari biosynth	n etic	organic hydroxy	aldehyde catabolic
response to granulocyte xenobiotic stimulus migration		response to wounding kinesis		response to	positive regulation chemokine-me	positive regulation of chemokine-mediated signaling pathway		of cardiac muscle contraction	vasculature development	intestinal sepithelial for structure	anatomical structure formation nvolved in rphogenesis	facial nerve structural organizatio	regulation of	sulfation sul	process heparin ation <sub>letabolic</sub>		catabolic proce	
lipid metabolic process	organic linole hydroxy acid compound metabolic process	d estrogen metabolic process	dermatan sulfate proteoglycan biosynthetic process	transduction by	netabolic process	organic acid netabolic process	oxalate metabolic process	rassalatars	muscle system process	maintenance regulation of metanephric nephron tubulecoa epithelial cell differentiation		organ sr senescence m	vein nooth uscle traction	3'-phosphoadenosine 5'-phosphosulfate metabolic process	proces homocysi metabolic p	teine ca	alcohol	catabolic process atecholamine metabolic process
phosphate-containing compound	regulation of protein phosphorylation	und phosphatidylethanolamin metabolic process ss	e phosphatidylethanolamine catabolic process	compound metabolic process	amine netabolic process	metabolic process	microtubule organization	intracellular signal transduction	negative regulation of transport	negative que regulation and of NF-kappaB import into	uaterna mmoniu group transpor	signalin	negative regulation of potassium ion transmembrane transporter activity	homeostasis	VCIS	ponse to	dendritic co	ell sulfur
metabolic process	nucleoside bisphosphate metabolic positive reg of proteas ubiquitin-de protein cat	regulation pendent of p <b>lipid</b> metabolic	regulation netabolism H3-K27	metabolic process	ohosphate   iosynthetic   process	phosphate	nistone	_	signal transductior	negative	hythmic	Tie	TORC2	regula no meostas of anion anatomical structure		imulus	differentiați dendritic co differentiati	
protein phosphorylation s	process unsatur small molecule fatty a biosynthetic metab	cid regulation o		biosynthetic process p	positive regulation of peptidyl–serine phosphorylation of STAT protein pyruvate	sphingolipi biosyntheti process	of protein metabolic process	negative regulation	negative negative regulation of signal	D-activating Of G-protein coupled	of signal egulation of cellular omponent	aling regulation of	regulation of cell	single-organism	n	motion	ingle-organisr process	<sup>n</sup> signaling
phosphorus metabolic process	steroid metabolic process MAF	process  cellular polysaccharide	regulation of microtubule-based process	molecule metabolic process astral microtubule organization	process  DP-N-acetylgalactosamine metabolic process	transfer activit fatty aci derivativ metabolic pr	y icosanoid metabolid process	of signaling  negative regulation of cell communication	transduction negative regulation of response to stimulus	pathway me	regulation of protein coupled eceptor protein gnaling pathway	transforming growth factor beta receptor complex assembly	ositive G-protein coupled of receptor logical signaling ocess pathway	cellula cellular componer component movement	nt <sub>communication</sub>	0	iological egulation	process  Ilti-organism process