## REVIGO Gene Ontology treemap

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regulation of nucleobase–containing compound metabolic process	regulatio <sup>g</sup> nitrogen con metabolic p	npound gene e	liation of I	egulation of piosynthetic process	cellular nitrogen compound biosynthet process	d biosyn	ound thetic	nucleobase–containing compound biosynthetic process	steroid catabolic process glutathione derivative	positive regulation cholester esterificat	n of nuclear-transcribe mRNA poly(A) tail shortening ion negative regulation of RNA	catabolic	negative regulation of cholesterol biosynthetic process small GTPase mediated	response to arsenic-containing substance negative regulation of		•	histone
aromatic compound biosynthetic process	RNA metabolio process	metabolic process	glutathion biosynthe process	tic metabol	ic from F s polymer prome	iption RNA compoi rase II	ease–containing ound metabolic process	positive regulation of metabolic process	metabolic process deoxycytidin metabolic	Ral ster	catabolic process old catabolic process decorpion of catabolic	cleoside positive regulation of deacetyles	transduction  Ras protein signal transduction	nistone H3–K2 acetylation response response to methyl methanesulfona	elongation e to arsenic— single organism reproductive	response to bacterial	ubstance
gene expression	glutathione derivative biosynthetic process negative	cellular nitrogen compound metabolic process	negative regulation of metaboli process cellular		from F polyme II prom	RNA (erase sile	gene encing	canonical Wnt signaling pathway involved in positive regulation of epithelial to mesenchymal transition	androgen catabolic process	Ral GTF activit deoxyribonuci metabolic pr	ty metabo proce lipopro	blic doxorubic metabol	cin pyrimidine	response to		detection of molecule of his	response to defenses of othe organism involver in symbiotic interaction
heterocycle biosynthetic process	regulation of macromolecule metabolic process positive regulation	formaldehyde biosynthetic positive r process mitochondrial	egulation of cellular metabolic	response to	protein <b>ynthesis</b> ficat	regulation	cellular macromole metaboli	cule	response to oxidative str	l to	JNK cascade	face morphoge	nesis moi	body rphogenesis	head development	chylomicron remnant clearance chylor	regulation of blood vessel endothelial cell migration
biosynthetic process	of cellular catabolic process organic cyclic compound	threonyl-tRNA aminoacylation peptidyl-proline		cGMP-mediated signaling	merapolic	stability autophagic cell death	hormor catabol proces	positive regulation of interleukin–6	toll-like receptor 6 signaling pathway	pathogen-associate molecular pattern dependent induction t symbiont of host innate immune respon	peripheral nervous system axon	retinal bipolar neu differentiat embryon	ic face mo	italia fate grphogenesis	association	remnant of regulation of store–operated calcium entry	
positive regulation of cellular	metabolic process cellular aromatic	hydroxylation to 3-hydroxy-L-proline positive regulatior of low-density	deoxyribonucleosid monophosphate biosynthetic proces	metabolic process	signaling pathway	positive regulation of seudopodium assembly	regulation of prote metabol process	on positive in regulation lic of cellular	protein kinase signaling cascade	premature senescence cellular	response to chemical stimulus cellular	hindgut	esis devel	ndgut ske	positive regulation of embryonic developments		nitrogen fixation
biosynthetic process regulation	compound metabolic process heterocycle	lipoprotein particle receptor catabolic process  positive regulation of transcription from RNA polymerase II	regulation of chromatir silencing regulation of sulfur	positive regulation of cytoplasmic mRNA processing body assembly regulation of type B	type B pancreatic cell apoptotic process  negative regulation of transcription	biosynthetic integration process signs	gulation of	transmission, cholinergic protein deubiquitination	response to hydroperoxide	alcohol	response to cellular response to ethanol	glucose–6–ph transpo ——— <b>gluc</b> protem in	osphate rt	secretory granule ocalization osphate tra	organophosphate ester transport nsport reverse	reproductive metal	abolic biologica
of metabolic process	metabolic process	promoter in response to calcium ion	metabolic process	pancreatic cell apoptotic process	elongation from RNA bi		tochondrial polarization	macromolecule metabolic process	RNA polymerase II	to heat	response to nitrosative stress	into nucle transloca	eus, p	hospholipid efflux	cholesterol transport		regulation